

SEQUENCE LISTING

<110> O'Brien, Timothy

<120> Repeat Sequences of the CA125 Gene and Their Use for Diagnostic and Therapeutic Interventions

<130> 40715-258841

<150> US 60/284,175

<151> 2001-04-17

<160> 306

<170> PatentIn version 3.0

<210> 1

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<213> Homo sapiens

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102300 825900

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Asp Thr Tyr Val Gly Pro Leu Tyr
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ggt ctg ctg agg cct gtg ttc aag aac acc agt gtt ggc cct ctg tac 96
Gly Leu Leu Arg Pro Val Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr
20 25 30

tct ggc tgc aga ctg acc ttg ctc agg ccc aag aag gat ggg gca gcc 144
 Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala Ala
 35 40 45

acc aaa gtg gat gcc atc tgc acc tac cgc cct gat ccc aaa agc cct 192
 Thr Lys Val Asp Ala Ile Cys Thr Tyr Arg Pro Asp Pro Lys Ser Pro
 50 55 60

gga ctg gac aga gag cag cta tac tgg gag ctg agc cag ggt gat gca 240
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<213> Homo sapiens

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Arg Arg Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln
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Gly Leu Leu Arg Pro Val Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr
 20 25 30

Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala Ala
 35 40 45

Thr Lys Val Asp Ala Ile Cys Thr Tyr Arg Pro Asp Pro Lys Ser Pro
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Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Gly Asp Ala
 65 70 75 80

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<212> DNA

<213> Artificial

<220>

<223> Synthetic Primer

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 Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu
 20 25 30
 Lys Asp Gly Thr Ala Thr Gly Val Asp Ala Ile Cys Thr His His Pro
 35 40 45
 Asp Pro Lys Ser Pro Arg Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu
 50 55 60
 Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly Pro Tyr Ala Leu Asp
 65 70 75 80
 Asn Asp Ser Leu Phe Val Asn Gly Phe Thr His Arg Ser Ser Val Ser
 85 90 95
 Thr Thr Ser Thr Pro Gly Thr Pro Thr Val Tyr Leu Gly Ala Ser Lys
 100 105 110
 Thr Pro Ala Ser Ile Phe Gly Pro Ser Ala Ala Ser Pro Leu Leu Ile
 115 120 125
 Pro Phe Thr
 130

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 <213> Homo sapiens

<400> 12
 Glu Arg Val Leu Gln Gly Leu Leu Met Pro Leu Phe Lys Asn Thr Ser
 1 5 10 15
 Val Ser Ser Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu
 20 25 30
 Lys Asp Gly Ala Ala Thr Arg Ala Asp Ala Val Cys Thr His Arg Pro

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          35              40              45
Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Lys Leu
 50              55              60

Ser Gln Leu Thr His Gly Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp
65              70              75              80

Arg His Ser Leu Tyr Val Asn Gly Phe Thr His Gln Ser Ser Met Thr
      85              90              95

Thr Thr Arg Thr Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser Arg
      100              105              110

Thr Pro Ala Ser Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu Ile
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Pro Phe
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<210> 13

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<212> PRT

<213> Homo sapiens

<400> 13

Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Ile Phe Lys Asn Thr Ser
1              5              10              15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Ser Glu
      20              25              30

Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Ile His Arg Leu
      35              40              45

Asp Pro Lys Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp Glu Leu
50              55              60

Ser Lys Leu Thr Asn Asp Ile Glu Glu Leu Gly Pro Tyr Thr Leu Asp
65              70              75              80

Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Gln Ser Ser Val Ser
      85              90              95

Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Arg Thr Ser Gly
      100              105              110

Thr Pro Ser Ser Leu Ser Ser Pro Thr Ile Met Ala Ala Gly Pro Leu
      115              120              125

Leu Ile Pro Phe
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<210> 14

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<212> PRT

<213> Homo sapiens

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Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Met Phe Lys Asn Thr Ser
1 5 10 15

Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu
20 25 30

Lys Asn Gly Ala Ala Thr Gly Met Asp Ala Ile Cys Ser His Arg Leu
35 40 45

Asp Pro Lys Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp Glu Leu
50 55 60

Ser Gln Leu Thr His Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp
65 70 75 80

Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Ala
85 90 95

Pro Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly
100 105 110

Thr Pro Ser Ser Leu Pro Ser Pro Thr Thr Ala Val Pro Leu Leu Ile
115 120 125

Pro Phe
130

<210> 15

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<212> PRT

<213> Homo sapiens

<400> 15

Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Leu Phe Lys Asn Ser Ser
1 5 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Ile Ser Leu Arg Ser Glu
20 25 30

Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His His Leu
35 40 45

Asn Pro Gln Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Gln Leu
50 55 60

Ser Gln Met Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp
65 70 75 80

Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Gly Leu
85 90 95

Thr Thr Ser Thr Pro Trp Thr Ser Thr Val Asp Leu Gly Thr Ser Gly
100 105 110

Thr Pro Ser Pro Val Pro Ser Pro Thr Thr Ala Gly Pro Phe Leu Ile
115 120 125

Pro Phe
130

<210> 16

<211> 130

<212> PRT

<213> Homo sapiens

<400> 16

Glu Arg Val Leu Gln Gly Leu Leu Arg Pro Leu Phe Lys Ser Thr Ser
1 5 10 15

Ala Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu
20 25 30

Lys His Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr Leu Arg Leu
35 40 45

Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu
50 55 60

Ser Gln Leu Thr Asn Ser Val Thr Glu Leu Gly Pro Tyr Thr Leu Asp
65 70 75 80

Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Pro
85 90 95

Thr Thr Ser Ile Pro Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly
100 105 110

Thr Pro Ala Ser Leu Pro Gly His Thr Ala Pro Gly Pro Leu Leu Ile
115 120 125

Pro Phe
130

<210> 17

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<213> Homo sapiens

<400> 17

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser
1 5 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu
20 25 30

Lys Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu
35 40 45

Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu
50 55 60

Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Thr Leu Asp
65 70 75 80

Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Pro
85 90 95

Thr Thr Ser Ile Pro Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly
100 105 110

Thr Pro Ala Ser Leu Pro Gly His Ile Val Pro Gly Pro Leu Leu Ile
115 120 125

Pro Phe
130

<210> 18

<211> 131

<212> PRT

<213> Homo sapiens

<400> 18

Glu Arg Val Leu Gln Gly Leu Leu Thr Pro Leu Phe Lys Asn Thr Ser
1 5 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu
20 25 30

Lys Gln Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Val
35 40 45

Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu
50 55 60

Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp
65 70 75 80

Arg Asp Ser Leu Tyr Val Asp Gly Phe Asn Pro Trp Ser Ser Val Pro
85 90 95

Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala Thr Ser Gly
100 105 110

Thr Pro Ser Pro Leu Pro Gly His Thr Ala Pro Val Pro Leu Leu Ile
115 120 125

Pro Phe Thr
130

<210> 19

<211> 131

<212> PRT

<213> Homo sapiens

<400> 19

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser
1 5 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu
20 25 30

Lys His Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu
35 40 45

Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu
50 55 60

Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp
65 70 75 80

Arg Gly Ser Leu Tyr Val Asn Gly Phe Thr His Arg Asn Phe Val Pro
85 90 95

Ile Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Gly Thr Ser Glu
100 105 110

Thr Pro Ser Ser Leu Pro Arg Pro Ile Val Pro Gly Pro Leu Leu Val
115 120 125

Pro Phe Thr
130

<210> 20

<211> 130

<212> PRT

<213> Homo sapiens

<400> 20

Glu Arg Val Leu Gln Gly Leu Leu Ser Pro Ile Phe Lys Asn Ser Ser
1 5 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu
20 25 30

Lys Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro
35 40 45

Asn Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu
50 55 60

Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp
65 70 75 80

Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Gln Asn Ser Val Pro
85 90 95

Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr Gly
100 105 110

Thr Pro Ser Ser Phe Pro Gly His Thr Glu Pro Gly Pro Leu Leu Ile
115 120 125

Pro Phe
130

<210> 21

<211> 131

<212> PRT

<213> Homo sapiens

<400> 21

Glu Arg Val Leu Gln Gly Leu Leu Arg Pro Leu Phe Lys Asn Thr Ser
1 5 10 15

Ile Gly Pro Leu Tyr Ser Ser Cys Arg Leu Thr Leu Leu Arg Pro Glu
20 25 30

Lys Asp Lys Ala Ala Thr Arg Val Asp Ala Ile Cys Thr His His Pro
35 40 45

Asp Pro Gln Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp Glu Leu
50 55 60

Ser Gln Leu Thr His Gly Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp
65 70 75 80

Arg Asp Ser Leu Tyr Val Asp Gly Phe Thr His Trp Ser Pro Ile Pro
85 90 95

Thr Thr Ser Thr Pro Gly Thr Ser Ile Val Asn Leu Gly Thr Ser Gly
100 105 110

Ile Pro Pro Ser Leu Pro Glu Thr Thr Ala Thr Gly Pro Leu Leu Ile
115 120 125

Pro Phe Thr
130

<210> 22

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<212> PRT

<213> Homo sapiens

<400> 22

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Arg Asn Ser Ser
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Leu Glu Tyr Leu Tyr Ser Gly Cys Arg Leu Ala Ser Leu Arg Pro Glu
20 25 30

Lys Asp Ser Ser Ala Met Ala Val Asp Ala Ile Cys Thr His Arg Pro
35 40 45

Asp Pro Glu Asp Leu Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu
50 55 60

Ser Asn Leu Thr Asn Gly Ile Gln Glu Leu Gly Pro Tyr Thr Leu Asp
65 70 75 80

Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Met Pro
85 90 95

Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Val Gly Thr Ser Gly
100 105 110

Thr Pro Ser Ser Ser Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Met
115 120 125

Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp
130 135 140

Met Arg Arg Thr Gly Ser Arg Lys Phe Asn Thr Met Glu Arg Val Leu

145 150 155 160
 Gln Gly Leu Leu Ser Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu
 165 170 175
 Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu Lys Asp Gly Ala
 180 185 190
 Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro Asn Pro Lys Arg
 195 200 205
 Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr
 210 215 220
 His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp Arg Asp Ser Leu
 225 230 235 240
 Tyr Val Asn Gly Phe Thr His Gln Asn Ser Val Pro Thr Thr Ser Thr
 245 250 255
 Pro Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr Gly Thr Pro Ser Ser
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 Phe Pro Gly His Thr Glu Pro Gly Pro Leu
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 <400> 23
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 Leu Glu Tyr Leu Tyr Ser Gly Cys Arg Leu Ala Ser Leu Arg Pro Glu
 20 25 30
 Lys Asp Ser Ser Ala Met Ala Val Asp Ala Ile Cys Thr His Arg Pro
 35 40 45
 Asp Pro Glu Asp Leu Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu
 50 55 60
 Ser Asn Leu Thr Asn Gly Ile Gln Glu Leu Gly Pro Tyr Thr Leu Asp
 65 70 75 80
 Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Gly Leu
 85 90 95
 Thr Thr Ser Thr Pro Trp Thr Ser Thr Val Asp Leu Gly Thr Ser Gly
 100 105 110

Thr Pro Ser Pro Val Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Ile
 115 120 125
 Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn
 130 135 140
 Met Gly His Pro Gly Ser Arg Lys Phe Asn Ile Met Glu Arg Val Leu
 145 150 155 160
 Gln Gly Leu Leu Met Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu
 165 170 175
 Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala
 180 185 190
 Ala Thr Arg Val Asp Ala Val Cys Thr Gln Arg Pro Asp Pro Lys Ser
 195 200 205
 Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Lys Leu Ser Gln Leu Thr
 210 215 220
 His Gly Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg His Ser Leu
 225 230 235 240
 Tyr Val Asn Gly Leu Thr His Gln Ser Ser Met Thr Thr Thr Arg Thr
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 Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser Arg Thr Pro Ala Ser
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 Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu Ile Pro Phe
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 Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser
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 Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu
 20 25 30
 Lys Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu
 35 40 45
 Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu
 50 55 60
 Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp
 65 70 75 80

Arg Gly Ser Leu Tyr Val Asn Gly Phe Thr His Arg Thr Ser Val Pro	85	90	95	
Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly	100	105	110	
Thr Pro Phe Ser Leu Pro Ser Pro Ala Thr Ala Gly Pro Leu Leu Val	115	120	125	
Leu Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Lys Tyr Glu Glu Asp	130	135	140	
Met His Arg Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu	145	150	155	160
Gln Thr Leu Leu Gly Pro Met Phe Lys Asn Thr Ser Val Gly Leu Leu	165	170	175	
Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Ser Glu Lys Asp Gly Ala	180	185	190	
Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser	195	200	205	
Pro Gly Val Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr	210	215	220	
Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu	225	230	235	240
Tyr Val Asn Gly Phe Thr His Trp Ile Pro	245	250		
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<211>	286			
<212>	PRT			
<213>	Homo sapiens			
<400>	25			
Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser	5	10	15	
Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu	20	25	30	
Lys Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu	35	40	45	
Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu	50	55	60	
Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asn				

65 70 75 80
 Arg Gly Ser Leu Tyr Val Asn Gly Phe Thr His Arg Asn Phe Val Pro
 85 90 95
 Ile Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Gly Thr Ser Glu
 100 105 110
 Thr Pro Ser Ser Leu Pro Arg Pro Ile Val Pro Gly Pro Leu Leu Ile
 115 120 125
 Pro Phe Thr Ile Asn Phe Thr Ile Thr Asn Leu Arg Tyr Glu Glu Asn
 130 135 140
 Met His His Pro Gly Ser Arg Lys Phe Asn Ile Met Glu Arg Val Leu
 145 150 155 160
 Gln Gly Leu Leu Gly Pro Leu Phe Lys Asn Ser Ser Val Gly Pro Leu
 165 170 175
 Tyr Ser Gly Cys Arg Leu Ile Ser Leu Arg Ser Glu Lys Asp Gly Ala
 180 185 190
 Ala Thr Gly Val Asp Ala Ile Cys Thr His His Leu Asn Pro Gln Ser
 195 200 205
 Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Gln Leu Ser Gln Met Thr
 210 215 220
 Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu
 225 230 235 240
 Tyr Val Asn Gly Phe Thr His Arg Ser Ser Gly Leu Thr Thr Ser Thr
 245 250 255
 Pro Trp Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Pro
 260 265 270
 Val Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Ile Pro Phe
 275 280 285
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 <400> 26
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 20 25 30

Lys Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu
 35 40 45
 Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu
 50 55 60
 Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp
 65 70 75 80
 Arg Gly Ser Leu Tyr Val Asn Gly Phe Ser Arg Gln Ser Ser Met Thr
 85 90 95
 Thr Thr Arg Thr Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser Arg
 100 105 110
 Thr Pro Ala Ser Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu Ile
 115 120 125
 Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn
 130 135 140
 Met Gly His Pro Gly Ser Arg Lys Phe Asn Ile Met Glu Arg Val Leu
 145 150 155 160
 Gln Gly Leu Leu Asn Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu
 165 170 175
 Tyr Ser Gly Cys Arg Leu Thr Ser Leu Lys Pro Glu Lys Asp Gly Ala
 180 185 190
 Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro Asn Pro Lys Arg
 195 200 205
 Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr
 210 215 220
 His Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu
 225 230 235 240
 Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Ala Pro Thr Ser Thr
 245 250 255
 Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Ser
 260 265 270
 Leu Pro Ser Pro Thr Thr Ala Val Pro Leu Leu Ile Pro Phe
 275 280 285

<210> 27

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<213> Homo sapiens

<400> 27

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser
 1 5 10 15
 Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu
 20 25 30
 Lys Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu
 35 40 45
 Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu
 50 55 60
 Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp
 65 70 75 80
 Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Pro
 85 90 95
 Thr Thr Ser Ile Pro Gly Thr Ser Ala Val His Leu Glu Thr Phe Gly
 100 105 110
 Thr Pro Ala Ser Leu His Gly His Thr Ala Pro Gly Pro Val Leu Val
 115 120 125
 Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp
 130 135 140
 Met Arg His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu
 145 150 155 160
 Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu
 165 170 175
 Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Arg Gly Ala
 180 185 190
 Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu Asp Pro Leu Asn
 195 200 205
 Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr
 210 215 220
 Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp Arg Gly Ser Leu
 225 230 235 240
 Tyr Val Asn Gly Phe Thr His Arg Asn Phe Val Pro Ile Thr Ser Thr
 245 250 255
 Pro Gly Thr Ser Thr Val His Leu Gly Thr Ser Glu Thr Pro Ser Ser
 260 265 270
 Leu Pro Arg Pro Ile Val Pro Gly Pro Leu Leu Ile Pro Phe
 275 280 285

<210> 28

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<213> Homo sapiens

<400> 28

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser
 1 5 10 15
 Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu
 20 25 30
 Lys His Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr Leu Arg Leu
 35 40 45
 Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu
 50 55 60
 Ser Gln Leu Thr Asn Ser Val Thr Glu Leu Gly Pro Tyr Thr Leu Asp
 65 70 75 80
 Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Pro
 85 90 95
 Thr Thr Ser Ile Pro Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly
 100 105 110
 Thr Pro Ala Ser Leu Pro Gly His Thr Ala Pro Gly Pro Leu Leu Val
 115 120 125
 Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp
 130 135 140
 Met Arg His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu
 145 150 155 160
 Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu
 165 170 175
 Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Arg Gly Ala
 180 185 190
 Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu Asp Pro Leu Asn
 195 200 205
 Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr
 210 215 220
 Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp Arg Gly Ser Leu
 225 230 235 240
 Tyr Val Asn Gly Phe Thr His Arg Asn Phe Val Pro Ile Thr Ser Thr
 245 250 255
 Pro Gly Thr Ser Thr Val His Leu Gly Thr Ser Glu Thr Pro Ser Ser
 260 265 270

Leu Pro Arg Pro Ile Val Pro Gly Pro Leu Leu Ile Pro Phe
 275 280 285

<210> 29

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<213> Homo sapiens

<400> 29

Glu Arg Val Leu Gln Gly Leu Leu Thr Pro Leu Phe Lys Asn Thr Ser
 1 5 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu
 20 25 30

Lys Gln Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Val
 35 40 45

Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu
 50 55 60

Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp
 65 70 75 80

Arg Asp Ser Leu Tyr Val Asn Gly Phe Asn Pro Trp Ser Ser Val Pro
 85 90 95

Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala Thr Ser Gly
 100 105 110

Thr Pro Ser Ser Leu Pro Gly His Thr Ala Pro Val Pro Leu Leu Ile
 115 120 125

Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu His Tyr Glu Glu Asn
 130 135 140

Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu
 145 150 155 160

Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu
 165 170 175

Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys His Gly Ala
 180 185 190

Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser
 195 200 205

Pro Gly Val Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr
 210 215 220

Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu
 225 230 235 240

Tyr Val Asn Gly Phe Thr His Trp Ile Pro Val Pro Thr Ser Ser Thr
245 250 255

Pro Gly Thr Ser Thr Val Asp Leu Gly Ser Gly Thr Pro Ser Ser Leu
260 265 270

Pro Ser Pro Thr Thr Ala Gly Pro Leu
275 280

<210> 30

<211> 217

<212> PRT

<213> Homo sapiens

<400> 30

Glu Arg Val Leu Gln Gly Leu Leu Arg Pro Leu Phe Lys Asn Thr Ser
1 5 10 15

Ile Gly Pro Leu Tyr Ser Ser Cys Arg Leu Thr Leu Leu Arg Pro Glu
20 25 30

Lys Asp Lys Ala Ala Thr Arg Val Asp Ala Ile Cys Thr His His Pro
35 40 45

Asp Pro Gln Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp Glu Leu
50 55 60

Ser Gln Leu Thr His Gly Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp
65 70 75 80

Arg Asp Ser Leu Tyr Val Asp Gly Phe Thr His Trp Ser Pro Ile Pro
85 90 95

Thr Thr Ser Thr Pro Gly Thr Ser Ile Val Asn Leu Gly Thr Ser Gly
100 105 110

Ile Pro Pro Ser Leu Pro Glu Thr Thr Ala Thr Gly Pro Leu Leu Ile
115 120 125

Pro Phe Thr Pro Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp
130 135 140

Met Arg Arg Thr Gly Ser Arg Lys Phe Asn Thr Met Glu Arg Val Leu
145 150 155 160

Gln Gly Leu Leu Ser Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu
165 170 175

Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu Lys Asp Gly Ala
180 185 190

Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro Asn Pro Lys Arg

195 200 205
 Pro Gly Leu Asp Arg Glu Gln Leu Tyr
 210 215

 <210> 31
 <211> 286
 <212> PRT
 <213> Homo sapiens

 <400> 31

 Glu Arg Val Leu Gln Gly Leu Leu Arg Pro Val Phe Lys Asn Thr Ser
 1 5 10 15
 Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys
 20 25 30
 Lys Asp Gly Ala Ala Thr Lys Val Asp Ala Ile Cys Thr Tyr Arg Pro
 35 40 45
 Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu
 50 55 60
 Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp
 65 70 75 80
 Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr Gln Arg Ser Ser Val Pro
 85 90 95
 Thr Thr Ser Ile Pro Gly Thr Pro Thr Val Asp Leu Gly Thr Ser Gly
 100 105 110
 Thr Pro Val Ser Lys Pro Gly Pro Ser Ala Ala Ser Pro Leu Leu Val
 115 120 125
 Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp
 130 135 140
 Met His Arg Pro Gly Ser Arg Lys Phe Asn Ala Thr Glu Arg Val Leu
 145 150 155 160
 Gln Gly Leu Leu Ser Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu
 165 170 175
 Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu Lys Asp Gly Ala
 180 185 190
 Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro Asn Pro Lys Arg
 195 200 205
 Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr
 210 215 220

His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp Arg Asp Ser Leu
225 230 235 240

Tyr Val Asn Gly Phe Thr His Gln Ser Ser Met Thr Thr Thr Arg Thr
245 250 255

Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser Arg Thr Pro Ala Ser
260 265 270

Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu Ile Pro Phe
275 280 285

<210> 32

<211> 288

<212> PRT

<213> Homo sapiens

<400> 32

Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Met Phe Lys Asn Thr Ser
1 5 10 15

Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys
20 25 30

Lys Asp Gly Ala Ala Thr Lys Val Asp Ala Ile Cys Thr Tyr Arg Pro
35 40 45

Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu
50 55 60

Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp
65 70 75 80

Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr Gln Arg Ser Ser Val Pro
85 90 95

Thr Thr Ser Ile Pro Gly Thr Pro Thr Val Asp Leu Gly Thr Ser Gly
100 105 110

Thr Pro Val Ser Lys Pro Gly Pro Ser Ala Ala Ser Pro Leu Leu Ile
115 120 125

Pro Phe Thr Ile Asn Phe Thr Ile Thr Asn Leu Arg Tyr Glu Glu Asn
130 135 140

Met Gly His Pro Gly Ser Arg Lys Phe Asn Ile Met Glu Arg Val Leu
145 150 155 160

Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu
165 170 175

Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala
180 185 190

Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser
195 200 205

Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr
210 220

Asn Asp Ile Glu Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu
225 230 235 240

Tyr Val Asn Gly Phe Thr His Gln Ser Ser Val Ser Thr Thr Ser Thr
245 250 255

Pro Gly Thr Ser Thr Val Asp Leu Arg Thr Ser Gly Thr Pro Ser Ser
260 265 270

Leu Ser Ser Pro Thr Ile Met Ala Ala Gly Pro Leu Leu Ile Pro Phe
275 280 285

<210> 33

<211> 284

<212> PRT

<213> Homo sapiens

<400> 33

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser
1 5 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu
20 25 30

Lys Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro
35 40 45

Asn Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu Tyr Cys Glu Leu
50 55 60

Ser Gln Leu Thr His Asp Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp
65 70 75 80

Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Gln Asn Ser Val Pro
85 90 95

Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr Gly
100 105 110

Thr Pro Ser Ser Phe Pro Gly His Thr Glu Pro Gly Pro Leu Leu Ile
115 120 125

Pro Phe Thr Phe Asn Phe Thr Ile Thr Asn Leu His Tyr Glu Glu Asn
130 135 140

Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu

145 150 155 160
 Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu
 165 170 175
 Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys His Glu Ala
 180 185 190
 Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Val Asp Pro Ile Gly
 195 200 205
 Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr
 210 215 220
 Asn Ser Ile His Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu
 225 230 235 240
 Tyr Val Asn Gly Phe Asn Pro Arg Ser Ser Val Pro Thr Thr Ser Thr
 245 250 255
 Pro Gly Thr Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Ser
 260 265 270
 Leu Pro Gly His Thr Ala Pro Val Pro Leu Leu Ile
 275 280
 <210> 34
 <211> 288
 <212> PRT
 <213> Homo sapiens

 <400> 34
 Glu Arg Val Leu Gln Gly Leu Leu Ser Pro Ile Ser Lys Asn Ser Ser
 1 5 10 15
 Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu
 20 25 30
 Lys Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro
 35 40 45
 Asn Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu
 50 55 60
 Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp
 65 70 75 80
 Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Gln Asn Ser Val Pro
 85 90 95
 Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr Gly
 100 105 110

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Thr Pro Ser Ser Phe Pro Gly His Thr Glu Pro Gly Pro Leu Leu Ile
115 120 125

Pro Phe Thr Val Asn Phe Thr Ile Thr Asn Leu Arg Tyr Glu Glu Asn
130 135 140

Met His His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu
145 150 155 160

Gln Gly Leu Leu Arg Pro Val Phe Lys Asn Thr Ser Val Gly Pro Leu
165 170 175

Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala
180 185 190

Ala Thr Lys Val Asp Ala Ile Cys Thr Tyr Arg Pro Asp Pro Lys Ser
195 200 205

Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr
210 215 220

Asn Asp Ile Glu Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu
225 230 235 240

Tyr Val Asn Gly Phe Thr His Gln Ser Ser Val Ser Thr Thr Ser Thr
245 250 255

Pro Gly Thr Ser Thr Val Asp Leu Arg Thr Ser Gly Thr Pro Ser Ser
260 265 270

Leu Ser Ser Pro Thr Ile Met Ala Ala Gly Pro Leu Leu Ile Pro Phe
275 280 285

<210> 35

<211> 274

<212> PRT

<213> Homo sapiens

<400> 35

Glu Arg Val Leu Gln Gly Leu Leu Ser Pro Ile Phe Lys Asn Ser Ser
1 5 10 15

Val Gly Ser Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu
20 25 30

Lys Asp Gly Ala Ala Thr Arg Val Asp Ala Val Cys Thr His Arg Pro
35 40 45

Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Lys Leu
50 55 60

Ser Gln Leu Thr His Gly Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp
65 70 75 80

Arg His Ser Leu Tyr Val Asn Gly Phe Thr His Gln Ser Ser Met Thr
85 90 95

Thr Thr Arg Thr Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser Arg
100 105 110

Thr Pro Ala Ser Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu Val
115 120 125

Leu Phe Thr Ile Asn Phe Thr Ile Thr Asn Gln Arg Tyr Glu Glu Asn
130 135 140

Met His His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu
145 150 155 160

Gln Gly Leu Leu Arg Pro Val Phe Lys Asn Thr Ser Val Gly Pro Leu
165 170 175

Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala
180 185 190

Ala Thr Lys Val Asp Ala Ile Cys Thr Tyr Arg Pro Asp Pro Lys Ser
195 200 205

Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr
210 215 220

His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Gln Asp Arg Asp Ser Leu
225 230 235 240

Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Pro Thr Thr Ser Ile
245 250 255

Pro Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly Thr Pro Ala Ser
260 265 270

Leu Pro

<210> 36

<211> 386

<212> PRT

<213> Homo sapiens

<400> 36

Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Met Phe Lys Asn Thr Ser
1 5 10 15

Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu
20 25 30

Lys Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu

35	40	45
Asp Pro Leu Asn Pro Gly Leu	Asp Arg Glu Gln Leu Tyr Trp Glu Leu	
50	55	60
Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp		
65	70	75
Arg Gly Ser Leu Tyr Val Asn Gly Phe Thr His Arg Asn Phe Val Pro		
85	90	95
Ile Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Gly Thr Ser Glu		
100	105	110
Thr Pro Ser Ser Leu Pro Arg Pro Ile Val Pro Gly Pro Leu Leu Val		
115	120	125
Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Ala		
130	135	140
Met Arg His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu		
145	150	155
Gln Gly Leu Leu Arg Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu		
165	170	175
Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala		
180	185	190
Ala Thr Arg Val Asp Ala Ala Cys Thr Tyr Arg Pro Asp Pro Lys Ser		
195	200	205
Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr		
210	215	220
His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Val Ser Leu		
225	230	235
Tyr Val Asn Gly Phe Asn Pro Arg Ser Ser Val Pro Thr Thr Ser Thr		
245	250	255
Pro Gly Thr Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Ser		
260	265	270
Leu Pro Gly His Thr Ala Pro Val Pro Leu Leu Ile Pro Phe Thr Leu		
275	280	285
Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro		
290	295	300
Gly Ser Arg Lys Phe Asn Thr Met Glu Arg Val Leu Gln Gly Leu Leu		
305	310	315
Arg Pro Leu Phe Lys Asn Thr Ser Ile Gly Pro Leu Tyr Ser Ser Cys		
325	330	335
Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Lys Ala Ala Thr Arg Val		
340	345	350

Asp Ala Ile Cys Thr His His Pro Asp Pro Gln Ser Pro Gly Leu Asn
355 360 365

Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His Gly Ile Thr
370 375 380

Glu Leu
385

<210> 37

<211> 438

<212> PRT

<213> Homo sapiens

<400> 37

Glu Arg Val Leu His Gly Leu Leu Thr Pro Leu Phe Lys Asn Thr Arg
1 5 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu
20 25 30

Lys Gln Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Val
35 40 45

Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu
50 55 60

Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp
65 70 75 80

Arg Asp Ser Leu Tyr Val Asn Gly Phe Asn Pro Trp Ser Ser Val Pro
85 90 95

Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala Thr Ser Gly
100 105 110

Thr Pro Ser Ser Leu Pro Gly His Thr Ala Pro Val Pro Leu Leu Ile
115 120 125

Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu His Tyr Glu Glu Asn
130 135 140

Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu
145 150 155 160

Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu
165 170 175

Tyr Ser Gly Cys Arg Leu Thr Leu Phe Lys Pro Glu Lys His Glu Ala
180 185 190

Ala Thr Gly Val Asp Ala Ile Cys Thr Leu Arg Leu Asp Pro Thr Gly
195 200 205

Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr
210 215 220

Asn Ser Val Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu
225 230 235 240

Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Pro Thr Thr Ser Ile
245 250 255

Pro Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly Thr Pro Ala Ser
260 265 270

Leu Pro Gly His Thr Ala Pro Gly Pro Leu Leu Ile Pro Phe Thr Leu
275 280 285

Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr
290 295 300

Gly Ser Arg Lys Phe Asn Thr Met Glu Arg Val Leu Gln Gly Leu Leu
305 310 315 320

Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys
325 330 335

Arg Leu Thr Leu Leu Arg Pro Glu Lys Arg Gly Ala Ala Thr Gly Val
340 345 350

Asp Thr Ile Cys Thr His Arg Leu Asp Pro Leu Asn Pro Gly Leu Asp
355 360 365

Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr Arg Gly Ile Ile
370 375 380

Glu Leu Gly Pro Tyr Leu Leu Asp Arg Gly Ser Leu Tyr Val Asn Gly
385 390 395 400

Phe Thr His Arg Asn Phe Val Pro Ile Thr Ser Thr Pro Gly Thr Ser
405 410 415

Thr Val His Leu Gly Thr Ser Glu Ile His Pro Ser Leu Pro Arg Pro
420 425 430

Ile Val Pro Gly Pro Leu
435

<210> 38

<211> 420

<212> PRT

<213> Homo sapiens

<400> 38

Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu Lys

1	5	10	15
Asp Gly Ala	Ala Thr Gly Met	Asp Ala Val Cys Leu Tyr	His Pro Asn
20		25	30
Pro Lys Arg	Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp	Glu Leu Ser	
35	40	45	
Gln Leu Thr His Asn Ile Thr	Glu Leu Gly Pro Tyr Ser Leu Asp Arg		
50	55	60	
Asp Ser Leu Tyr Val Asn Gly Phe Thr His Gln Asn Ser Val Pro Thr			
65	70	75	80
Thr Ser Thr Pro Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr Gly Thr			
85	90	95	
Pro Ser Ser Phe Pro Gly His Thr Glu Pro Gly Pro Leu Leu Ile Pro			
100	105	110	
Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn Met			
115	120	125	
Gly His Pro Gly Ser Arg Lys Phe Asn Ile Thr Glu Ser Val Leu Gln			
130	135	140	
Gly Leu Leu Thr Pro Leu Phe Lys Asn Ser Ser Val Gly Pro Leu Tyr			
145	150	155	160
Ser Gly Cys Arg Leu Ile Ser Leu Arg Ser Glu Lys Asp Gly Ala Ala			
165	170	175	
Thr Gly Val Asp Ala Ile Cys Thr His His Leu Asn Pro Gln Ser Pro			
180	185	190	
Gly Leu Asp Arg Glu Gln Leu Tyr Trp Gln Leu Ser Gln Met Thr Asn			
195	200	205	
Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr			
210	215	220	
Val Asn Gly Phe Thr His Arg Ser Leu Gly Leu Thr Thr Ser Thr Pro			
225	230	235	240
Trp Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Pro Val			
245	250	255	
Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Ile Pro Phe Thr Leu Asn			
260	265	270	
Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn Met Gly His Pro Gly			
275	280	285	
Ser Arg Lys Phe Asn Ile Met Glu Arg Val Leu Gln Gly Leu Leu Arg			
290	295	300	
Pro Val Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg			
305	310	315	320

Ala	Thr	Leu	Leu	Arg	Pro	Lys	Lys	Asp	Gly	Ala	Ala	Thr	Lys	Val	Asp
				325					330					335	
Ala	Ile	Cys	Thr	Tyr	Arg	Pro	Asp	Pro	Lys	Ser	Pro	Gly	Leu	Asp	Arg
				340				345					350		
Glu	Gln	Leu	Tyr	Trp	Glu	Leu	Ser	Gln	Leu	Thr	His	Ser	Ile	Thr	Glu
		355					360						365		
Leu	Gly	Pro	Tyr	Thr	Leu	Asp	Arg	Asp	Ser	Leu	Tyr	Val	Asn	Gly	Phe
						375					380				
Thr	Gln	Arg	Ser	Ser	Val	Pro	Thr	Thr	Ser	Ile	Pro	Gly	Thr	Pro	Thr
385					390					395				400	
Val	Asp	Leu	Gly	Thr	Ser	Gly	Thr	Pro	Val	Ser	Lys	Pro	Gly	Pro	Ser
				405					410					415	
Ala	Ala	Ser	Pro												
			420												
<210> 39															
<211> 439															
<212> PRT															
<213> Homo sapiens															
<400> 39															
Glu	Arg	Val	Leu	Gln	Gly	Pro	Leu	Ser	Pro	Ile	Phe	Lys	Asn	Ser	Ser
1			5						10					15	
Val	Gly	Pro	Leu	Tyr	Ser	Gly	Cys	Arg	Leu	Thr	Ser	Leu	Arg	Pro	Glu
			20					25					30		
Lys	Asp	Gly	Ala	Ala	Thr	Gly	Met	Asp	Ala	Val	Cys	Leu	Tyr	His	Pro
		35					40					45			
Asn	Pro	Lys	Arg	Pro	Gly	Leu	Asp	Arg	Glu	Gln	Leu	Tyr	Trp	Glu	Leu
	50					55					60				
Ser	Gln	Leu	Thr	His	Asn	Ile	Thr	Glu	Leu	Gly	Pro	Tyr	Ser	Leu	Asp
65					70					75				80	
Arg	Asp	Ser	Leu	Tyr	Val	Asn	Gly	Phe	Thr	His	Gln	Asn	Ser	Val	Pro
				85					90					95	
Thr	Thr	Ser	Thr	Pro	Gly	Thr	Ser	Thr	Val	Tyr	Trp	Ala	Thr	Thr	Gly
			100					105					110		
Thr	Pro	Ser	Ser	Phe	Pro	Gly	His	Thr	Glu	Pro	Gly	Pro	Leu	Leu	Ile
			115				120					125			
Pro	Phe	Thr	Leu	Asn	Phe	Thr	Ile	Thr	Asn	Leu	Gln	Tyr	Glu	Glu	Asn
	130					135					140				

Met Gly His Pro Gly Ser Arg Lys Phe Asn Ile Thr Glu Arg Val Leu
 145 150 155 160
 Gln Gly Leu Leu Asn Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu
 165 170 175
 Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu Lys Asp Gly Ala
 180 185 190
 Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro Asn Pro Lys Arg
 195 200 205
 Pro Gly Leu Asp Arg Glu Gln Leu Tyr Cys Glu Leu Ser Gln Leu Thr
 210 215 220
 His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp Arg Asp Ser Leu
 225 230 235
 Tyr Val Asn Gly Phe Thr His Gln Asn Ser Val Pro Thr Thr Ser Thr
 245 250 255
 Pro Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr Gly Thr Pro Ser Ser
 260 265 270
 Phe Pro Gly His Thr Glu Pro Gly Pro Leu Leu Ile Pro Phe Thr Leu
 275 280 285
 Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr
 290 295 300
 Gly Ser Arg Lys Phe Asn Thr Met Glu Arg Val Leu Gln Gly Leu Leu
 305 310 315 320
 Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys
 325 330 335
 Arg Leu Thr Leu Leu Arg Pro Glu Lys His Gly Ala Ala Thr Gly Val
 340 345 350
 Asp Ala Ile Cys Thr Leu Arg Leu Asp Pro Thr Gly Pro Gly Leu Asp
 355 360 365
 Arg Glu Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Ser Val Thr
 370 375 380
 Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly
 385 390 395 400
 Phe Thr His Arg Ser Ser Val Pro Thr Thr Ser Ile Pro Gly Thr Ser
 405 410 415
 Ala Val His Leu Glu Thr Ser Gly Thr Pro Ala Ser Leu Pro Gly His
 420 425 430
 Thr Ala Pro Gly Pro Leu Leu
 435

<210> 40

<211> 424

<212> PRT

<213> Homo sapiens

<400> 40

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Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg
1      5      10      15

Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Tyr Thr His
      20      25      30

Arg Leu Asp Pro Lys Ser Pro Gly Val Asp Arg Glu Gln Leu Tyr Trp
      35      40      45

Glu Leu Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr
      50      55      60

Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Gln Thr Ser
      65      70      75      80

Ala Pro Asn Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Thr
      85      90      95

Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr Ser Ala Gly Pro Leu
      100      105      110

Leu Ile Pro Phe Thr Ile Asn Phe Thr Ile Thr Asn Leu Arg Tyr Glu
      115      120      125

Glu Asn Met His His Pro Gly Ser Arg Lys Phe Asn Thr Met Glu Arg
      130      135      140

Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly
      145      150      155      160

Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp
      165      170      175

Gly Val Ala Thr Arg Val Asp Ala Ile Cys Thr His Arg Pro Asp Pro
      180      185      190

Lys Ile Pro Gly Leu Asp Arg Gln Gln Leu Tyr Trp Glu Leu Ser Gln
      195      200      205

Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp
      210      215      220

Ser Leu Tyr Val Asn Gly Phe Thr Gln Arg Ser Ser Val Pro Thr Thr
      225      230      235      240

Ser Thr Pro Gly Thr Phe Thr Val Gln Pro Glu Thr Ser Glu Thr Pro
      245      250      255

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Ser Ser Leu Pro Gly Pro Thr Ala Thr Gly Pro Val Leu Leu Pro Phe
260 265 270

Thr Leu Asn Phe Thr Ile Ile Asn Leu Gln Tyr Glu Glu Asp Met His
275 280 285

Arg Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly
290 295 300

Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser
305 310 315 320

Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys His Gly Ala Ala Thr
325 330 335

Gly Val Asp Ala Ile Cys Thr Leu Arg Leu Asp Pro Thr Gly Pro Gly
340 345 350

Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Ser
355 360 365

Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val
370 375 380

Asn Gly Phe Asn Pro Trp Ser Ser Val Pro Thr Thr Ser Thr Pro Gly
385 390 395 400

Thr Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Ser Leu Pro
405 410 415

Gly His Thr Ala Pro Val Pro Leu
420

<210> 41

<211> 418

<212> PRT

<213> Homo sapiens

<400> 41

Thr Leu Leu Arg Pro Lys Lys Asp Gly Val Ala Thr Gly Val Asp Ala
1 5 10 15

Ile Cys Thr His Arg Leu Asp Pro Lys Ser Pro Gly Leu Asn Arg Glu
20 25 30

Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr Asn Asp Ile Glu Glu Leu
35 40 45

Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr
50 55 60

His Gln Ser Ser Val Ser Thr Thr Ser Thr Pro Gly Thr Ser Thr Val

65					70					75					80
Asp	Leu	Arg	Thr	Ser	Gly	Thr	Pro	Ser	Ser	Leu	Ser	Ser	Pro	Thr	Ile
				85					90					95	
Met	Ala	Ala	Gly	Pro	Leu	Leu	Ile	Pro	Phe	Thr	Ile	Asn	Phe	Thr	Ile
			100					105					110		
Thr	Asn	Leu	Arg	Tyr	Glu	Glu	Asn	Met	His	His	Pro	Gly	Ser	Arg	Lys
		115					120					125			
Phe	Asn	Thr	Met	Glu	Arg	Val	Leu	Gln	Gly	Leu	Leu	Met	Pro	Leu	Phe
		130				135					140				
Lys	Asn	Thr	Ser	Val	Ser	Ser	Leu	Tyr	Ser	Gly	Cys	Arg	Leu	Thr	Leu
145					150					155					160
Leu	Arg	Pro	Glu	Lys	Asp	Gly	Ala	Ala	Thr	Arg	Val	Asp	Ala	Val	Cys
				165					170					175	
Thr	His	Arg	Pro	Asp	Pro	Lys	Ser	Pro	Gly	Leu	Asp	Arg	Glu	Arg	Leu
			180					185					190		
Tyr	Trp	Lys	Leu	Ser	Gln	Leu	Thr	His	Gly	Ile	Thr	Glu	Leu	Gly	Pro
		195					200					205			
Tyr	Thr	Leu	Asp	Arg	Asn	Ser	Leu	Tyr	Val	Asn	Gly	Phe	Thr	His	Arg
210						215					220				
Ser	Ser	Met	Pro	Thr	Thr	Ser	Thr	Pro	Gly	Thr	Ser	Thr	Val	Asp	Val
225					230					235					240
Gly	Thr	Ser	Gly	Thr	Pro	Ser	Ser	Ser	Pro	Ser	Pro	Thr	Thr	Ala	Gly
				245					250					255	
Pro	Leu	Leu	Met	Pro	Phe	Thr	Leu	Asn	Phe	Thr	Ile	Thr	Asn	Leu	Gln
			260					265					270		
Tyr	Glu	Glu	Asp	Met	Arg	Arg	Thr	Gly	Ser	Arg	Lys	Phe	Asn	Thr	Met
		275					280					285			
Glu	Arg	Val	Leu	Gln	Gly	Leu	Leu	Lys	Pro	Leu	Phe	Lys	Ser	Thr	Ser
290						295					300				
Val	Gly	Pro	Leu	Tyr	Ser	Gly	Cys	Arg	Leu	Thr	Leu	Leu	Arg	Pro	Glu
305					310					315					320
Lys	His	Gly	Ala	Ala	Thr	Gly	Val	Asp	Ala	Ile	Cys	Thr	Leu	Arg	Leu
			325						330					335	
Asp	Pro	Thr	Gly	Pro	Gly	Leu	Asp	Arg	Glu	Arg	Leu	Tyr	Trp	Glu	Leu
			340				345						350		
Ser	Gln	Leu	Thr	Asn	Ser	Val	Thr	Glu	Leu	Gly	Pro	Tyr	Thr	Leu	Asp
		355					360					365			
Arg	Asp	Ser	Leu	Tyr	Val	Asn	Gly	Phe	Thr	His	Arg	Ser	Ser	Val	Pro
370						375					380				

Thr Thr Ser Ile Pro Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly
385 390 395 400

Thr Pro Ala Ser Leu Pro Gly His Thr Ala Pro Gly Pro Leu Leu Ile
405 410 415

Pro Phe

<210> 42

<211> 443

<212> PRT

<213> Homo sapiens

<400> 42

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser
1 5 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu
20 25 30

Lys Asp Gly Val Ala Thr Arg Val Asp Ala Ile Cys Thr His Arg Pro
35 40 45

Asp Pro Lys Ile Pro Gly Leu Asp Arg Gln Gln Leu Tyr Trp Glu Leu
50 55 60

Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp
65 70 75 80

Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr Gln Arg Ser Ser Val Pro
85 90 95

Thr Thr Ser Thr Pro Gly Thr Phe Thr Val Gln Pro Glu Thr Ser Glu
100 105 110

Thr Pro Ser Ser Leu Pro Gly Pro Thr Ala Thr Gly Pro Val Leu Leu
115 120 125

Pro Phe Thr Leu Asn Phe Thr Ile Ile Asn Leu Gln Tyr Glu Glu Asp
130 135 140

Met His Arg Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu
145 150 155 160

Gln Gly Leu Leu Met Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu
165 170 175

Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Gln Glu Ala
180 185 190

Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu Asp Pro Ser Glu
195 200 205

Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr
 210 215 220
 Asn Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu
 225 230 235 240
 Tyr Val Asn Gly Phe Thr His Ser Gly Val Leu Cys Pro Pro Pro Ser
 245 250 255
 Ile Leu Gly Ile Phe Thr Val Gln Pro Glu Thr Phe Glu Thr Pro Ser
 260 265 270
 Ser Leu Pro Gly Pro Thr Ala Thr Gly Pro Val Leu Leu Pro Phe Thr
 275 280 285
 Leu Asn Phe Thr Ile Ile Asn Leu Gln Tyr Glu Glu Asp Met His Arg
 290 295 300
 Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu
 305 310 315 320
 Leu Met Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly
 325 330 335
 Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Gln Glu Ala Ala Thr Gly
 340 345 350
 Val Asp Thr Ile Cys Thr His Arg Val Asp Pro Ile Gly Pro Gly Leu
 355 360 365
 Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Ser Ile
 370 375 380
 Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn
 385 390 395 400
 Gly Phe Asn Pro Trp Ser Ser Val Pro Thr Thr Ser Thr Pro Gly Thr
 405 410 415
 Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Ser Leu Pro Gly
 420 425 430
 His Thr Ala Pro Val Pro Leu Leu Ile Pro Phe
 435 440
 <210> 43
 <211> 442
 <212> PRT
 <213> Homo sapiens
 <400> 43
 Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Arg Asn Ser Ser

1	5	10	15
Leu Glu Tyr	Leu Tyr Ser Gly	Cys Arg Leu Ala Ser Leu Arg	Pro Glu
	20	25	30
Lys Asp Ser	Ser Ala Met Ala Val Asp	Ala Ile Cys Thr	His Arg Pro
	35	40	45
Asp Pro Glu	Asp Leu Gly Leu Asp Arg	Glu Arg Leu Tyr Trp	Glu Leu
	50	55	60
Ser Asn Leu Thr	Asn Gly Ile Gln Glu	Leu Gly Pro Tyr Thr	Leu Asp
	65	70	75
Arg Asn Ser Leu	Tyr Val Asn Gly Phe Thr	His Arg Ser Ser Met	Pro
	85	90	95
Thr Thr Ser Thr	Pro Gly Thr Ser Thr	Val Asp Val Gly Thr	Ser Gly
	100	105	110
Thr Pro Ser Ser	Ser Pro Ser Pro Thr	Thr Ala Gly Pro	Leu Leu Met
	115	120	125
Pro Phe Thr Leu	Asn Phe Thr Ile Thr	Asn Leu Gln Tyr	Glu Glu Asp
	130	135	140
Met Arg Arg Thr	Gly Ser Arg Lys Phe Asn Thr	Met Glu Ser Val	Leu
	145	150	155
Gln Gly Leu Leu	Lys Pro Leu Phe Lys Asn Thr	Ser Val Gly Pro	Leu
	165	170	175
Tyr Ser Gly Cys	Arg Leu Thr Leu Leu Arg	Pro Lys Lys Asp	Gly Ala
	180	185	190
Ala Thr Gly Val	Asp Ala Ile Cys Thr	His Arg Leu Asp	Pro Lys Ser
	195	200	205
Pro Gly Leu Asn	Arg Glu Gln Leu Tyr Trp	Glu Leu Ser Lys	Leu Thr
	210	215	220
Asn Asp Ile Glu	Glu Val Gly Pro Tyr Thr	Leu Asp Arg Asn	Ser Leu
	225	230	235
Tyr Val Asn Gly	Phe Thr His Arg Ser Phe	Val Ala Pro Thr	Ser Thr
	245	250	255
Leu Gly Thr Ser	Thr Val Asp Leu Gly Thr	Ser Gly Thr Pro	Ser Ser
	260	265	270
Leu Pro Ser Pro	Thr Thr Gly Val Pro	Leu Leu Ile Pro	Phe Thr Leu
	275	280	285
Asn Phe Thr Ile	Thr Asn Leu Gln Tyr Glu	Glu Asn Met Gly	His Pro
	290	295	300
Gly Ser Arg Lys	Phe Asn Ile Met Glu Arg	Val Leu Gln Gly	Leu Leu
	305	310	315
			320

Met Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu Tyr Ser Gly Cys
 325 330 335
 Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala Ala Thr Arg Val
 340 345 350
 Val Ala Val Cys Thr His Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp
 355 360 365
 Arg Glu Arg Leu Tyr Trp Lys Leu Ser Gln Leu Thr His Gly Ile Thr
 370 375 380
 Glu Leu Gly Pro Tyr Thr Leu Asp Arg His Ser Leu Tyr Val Asn Gly
 385 390 395 400
 Phe Thr His Gln Ser Ser Met Thr Thr Thr Arg Thr Pro Asp Thr Ser
 405 410 415
 Thr Met His Leu Ala Thr Ser Arg Thr Pro Ala Ser Leu Ser Gly Pro
 420 425 430
 Thr Thr Ala Ser Pro Leu Leu Ile Pro Phe
 435 440
 <210> 44
 <211> 442
 <212> PRT
 <213> Homo sapiens
 <400> 44
 Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser
 1 5 10 15
 Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu
 20 25 30
 Lys Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu
 35 40 45
 Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu
 50 55 60
 Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp
 65 70 75 80
 Arg Gly Ser Leu Tyr Val Asn Gly Phe Thr His Arg Asn Phe Val Pro
 85 90 95
 Ile Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Gly Thr Ser Glu
 100 105 110
 Thr Pro Ser Ser Leu Pro Arg Pro Ile Val Pro Gly Pro Leu Leu Ile
 115 120 125

Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn
 130 135 140
 Met Gly His Pro Gly Ser Arg Lys Phe Asn Ile Thr Glu Arg Val Leu
 145 150 155 160
 Gln Gly Leu Leu Lys Pro Leu Phe Arg Asn Ser Ser Leu Glu Tyr Leu
 165 170 175
 Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu Lys Asp Ser Ser
 180 185 190
 Thr Met Ala Val Asp Ala Ile Cys Thr His Arg Pro Asp Pro Glu Asp
 195 200 205
 Leu Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Asn Leu Thr
 210 215 220
 Asn Gly Ile Gln Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu
 225 230 235 240
 Tyr Val Asn Gly Phe Thr His Arg Ser Phe Met Pro Thr Thr Ser Thr
 245 250 255
 Leu Gly Thr Ser Thr Val Asp Val Gly Thr Ser Gly Thr Pro Ser Ser
 260 265 270
 Ser Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Met Pro Phe Thr Leu
 275 280 285
 Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr
 290 295 300
 Gly Ser Arg Lys Phe Asn Thr Met Glu Ser Val Leu Gln Gly Leu Leu
 305 310 315 320
 Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys
 325 330 335
 Arg Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala Ala Thr Gly Val
 340 345 350
 Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser Pro Gly Leu Asn
 355 360 365
 Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr Asn Asp Ile Glu
 370 375 380
 Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly
 385 390 395 400
 Phe Thr His Gln Ser Ser Val Ser Thr Thr Ser Thr Pro Gly Thr Ser
 405 410 415
 Thr Val Asp Pro Arg Thr Ser Gly Thr Pro Ser Ser Leu Ser Ser Pro
 420 425 430
 Thr Ile Met Ala Ala Gly Pro Leu Leu Ile

435 440

<210> 45

<211> 379

<212> PRT

<213> Homo sapiens

<400> 45

Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Met Phe Lys Asn Thr Ser
 1 5 10 15

Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu
 20 25 30

Lys Asn Gly Ala Ala Thr Gly Met Asp Ala Ile Cys Ser His Arg Leu
 35 40 45

Asp Pro Lys Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp Glu Leu
 50 55 60

Ser Gln Leu Thr His Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp
 65 70 75 80

Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Ala
 85 90 95

Pro Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly
 100 105 110

Thr Pro Ser Ser Leu Pro Ser Pro Thr Thr Ala Val Pro Leu Leu Ile
 115 120 125

Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Lys Tyr Glu Glu Asp
 130 135 140

Met His Cys Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu
 145 150 155 160

Gln Ser Leu Phe Gly Pro Met Phe Lys Asn Thr Ser Val Gly Pro Leu
 165 170 175

Tyr Ser Gly Cys Arg Leu Thr Leu Phe Arg Ser Glu Lys Asp Gly Ala
 180 185 190

Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser
 195 200 205

Pro Gly Val Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr
 210 215 220

Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu
 225 230 235 240

Tyr Val Asn Gly Phe Thr His Gln Thr Ser Ala Pro Asn Thr Ser Thr
245 250 255

Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Ser
260 265 270

Leu Pro Ser Pro Thr Ser Ala Gly Pro Leu Leu Val Pro Phe Thr Leu
275 280 285

Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr
290 295 300

Gly Ser Arg Lys Phe Asn Thr Met Glu Ser Val Leu Gln Gly Leu Leu
305 310 315 320

Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys
325 330 335

Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala Ala Thr Gly Val
340 345 350

Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser Pro Gly Leu Asn
355 360 365

Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu
370 375

<210> 46

<211> 439

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (1)..(439)

<223> Any "X" = any amino acid

<400> 46

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser
1 5 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu
20 25 30

Lys His Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr Leu Arg Leu
35 40 45

Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu

50 55 60
 Ser Gln Leu Thr Asn Ser Val Thr Glu Leu Gly Pro Tyr Thr Leu Asp
 65 70 75 80
 Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Pro
 85 90 95
 Thr Thr Ser Ile Pro Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly
 100 105 110
 Thr Pro Ala Ser Leu Pro Gly His Thr Ala Pro Gly Pro Leu Leu Ile
 115 120 125
 Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu His Tyr Glu Glu Asn
 130 135 140
 Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Met Glu Arg Val Leu
 145 150 155 160
 Gln Gly Cys Leu Val Pro Cys Ser Arg Asn Thr Asn Val Gly Leu Leu
 165 170 175
 Tyr Ser Gly Cys Arg Leu Thr Leu Leu Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 180 185 190
 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 195 200 205
 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 210 215 220
 Xaa Xaa Xaa Xaa Xaa Xaa Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu
 225 230 235 240
 Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Ala Pro Thr Ser Thr
 245 250 255
 Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Ser
 260 265 270
 Leu Pro Ser Pro Thr Thr Val Pro Leu Leu Val Pro Phe Thr Leu Asn
 275 280 285
 Phe Thr Ile Thr Asn Leu Gln Tyr Gly Glu Asp Met Arg His Pro Gly
 290 295 300
 Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Gly
 305 310 315 320
 Pro Leu Phe Lys Asn Ser Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg
 325 330 335
 Leu Ile Ser Leu Arg Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp
 340 345 350
 Ala Ile Cys Thr His His Leu Asn Pro Gln Ser Pro Gly Leu Asp Arg
 355 360 365

Glu Gln Leu Tyr Trp Gln Leu Ser Gln Val Thr Asn Gly Ile Lys Glu
370 375 380

Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe
385 390 395 400

Thr His Arg Ser Ser Gly Leu Thr Thr Ser Thr Pro Trp Thr Ser Thr
405 410 415

Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Pro Val Pro Ser Pro Thr
420 425 430

Thr Ala Gly Pro Leu Leu Ile
435

<210> 47

<211> 1366

<212> PRT

<213> Homo sapiens

<400> 47

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Arg Asn Ser Ser
1 5 10 15

Leu Glu Tyr Leu Tyr Ser Gly Cys Arg Leu Ala Ser Leu Arg Pro Glu
20 25 30

Lys Asp Ser Ser Ala Met Ala Val Asp Ala Ile Cys Thr His Arg Pro
35 40 45

Asp Pro Glu Asp Leu Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu
50 55 60

Ser Asn Leu Thr Asn Gly Ile Gln Glu Leu Gly Pro Tyr Thr Leu Asp
65 70 75 80

Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Met Pro
85 90 95

Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Val Gly Thr Ser Gly
100 105 110

Thr Pro Ser Ser Ser Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Met
115 120 125

Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp
130 135 140

Met Arg Arg Thr Gly Ser Arg Lys Phe Asn Thr Met Glu Arg Val Leu
145 150 155 160

Gln Gly Pro Leu Ser Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu
165 170 175

Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu Lys Asp Gly Ala
 180 185 190
 Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro Asn Pro Lys Arg
 195 200 205
 Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr
 210 215 220
 His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp Arg Asp Ser Leu
 225 230 235 240
 Tyr Val Asn Gly Phe Thr His Gln Asn Ser Val Pro Thr Thr Ser Thr
 245 250 255
 Pro Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr Gly Thr Pro Ser Ser
 260 265 270
 Phe Pro Gly His Thr Glu Pro Gly Pro Leu Leu Ile Pro Phe Thr Leu
 275 280 285
 Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn Met Gly His Pro
 290 295 300
 Gly Ser Arg Lys Phe Asn Ile Thr Glu Arg Val Leu Gln Gly Leu Leu
 305 310 315 320
 Asn Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu Tyr Ser Gly Cys
 325 330 335
 Arg Leu Thr Ser Leu Arg Pro Glu Lys Asp Gly Ala Ala Thr Gly Met
 340 345 350
 Asp Ala Val Cys Leu Tyr His Pro Asn Pro Lys Arg Pro Gly Leu Asp
 355 360 365
 Arg Glu Gln Leu Tyr Cys Glu Leu Ser Gln Leu Thr His Asn Ile Thr
 370 375 380
 Glu Leu Gly Pro Tyr Ser Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly
 385 390 395 400
 Phe Thr His Gln Asn Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser
 405 410 415
 Thr Val Tyr Trp Ala Thr Thr Gly Thr Pro Ser Ser Phe Pro Gly His
 420 425 430
 Thr Glu Pro Gly Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile
 435 440 445
 Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr Gly Ser Arg Lys
 450 455 460
 Phe Asn Thr Met Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe
 465 470 475 480
 Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu

485										490					495				
Leu	Arg	Pro	Glu	Lys	His	Gly	Ala	Ala	Thr	Gly	Val	Asp	Ala	Ile	Cys				
			500						505						510				
Thr	Leu	Arg	Leu	Asp	Pro	Thr	Gly	Pro	Gly	Leu	Asp	Arg	Glu	Arg	Leu				
			515						520						525				
Tyr	Trp	Glu	Leu	Ser	Gln	Leu	Thr	Asn	Ser	Val	Thr	Glu	Leu	Gly	Pro				
			530						535						540				
Tyr	Thr	Leu	Asp	Arg	Asp	Ser	Leu	Tyr	Val	Asn	Gly	Phe	Thr	His	Arg				
			545						550						560				
Ser	Ser	Val	Pro	Thr	Thr	Ser	Ile	Pro	Gly	Thr	Ser	Ala	Val	His	Leu				
									565						575				
Glu	Thr	Ser	Gly	Thr	Pro	Ala	Ser	Leu	Pro	Gly	His	Thr	Ala	Pro	Gly				
			580						585						590				
Pro	Leu	Leu	Val	Pro	Phe	Thr	Leu	Asn	Phe	Thr	Ile	Thr	Asn	Leu	Gln				
			595						600						605				
Tyr	Glu	Glu	Asp	Met	Arg	His	Pro	Gly	Ser	Arg	Lys	Phe	Asn	Thr	Thr				
			610						615						620				
Glu	Arg	Val	Leu	Gln	Gly	Leu	Leu	Lys	Pro	Leu	Phe	Lys	Ser	Thr	Ser				
			625						630						640				
Val	Gly	Pro	Leu	Tyr	Ser	Gly	Cys	Arg	Leu	Thr	Leu	Leu	Arg	Pro	Glu				
									645						655				
Lys	Arg	Gly	Ala	Ala	Thr	Gly	Val	Asp	Thr	Ile	Cys	Thr	His	Arg	Leu				
									660						670				
Asp	Pro	Leu	Asn	Pro	Gly	Leu	Asp	Arg	Glu	Gln	Leu	Tyr	Trp	Glu	Leu				
									675						685				
Ser	Lys	Leu	Thr	Arg	Gly	Ile	Ile	Glu	Leu	Gly	Pro	Tyr	Leu	Leu	Asp				
									690						700				
Arg	Gly	Ser	Leu	Tyr	Val	Asn	Gly	Phe	Thr	His	Arg	Asn	Phe	Val	Pro				
									705						720				
Ile	Thr	Ser	Thr	Pro	Gly	Thr	Ser	Thr	Val	His	Leu	Gly	Thr	Ser	Glu				
									725						735				
Thr	Pro	Ser	Ser	Leu	Pro	Arg	Pro	Ile	Val	Pro	Gly	Pro	Leu	Leu	Ile				
									740						750				
Pro	Phe	Thr	Leu	Asn	Phe	Thr	Ile	Thr	Asn	Leu	Gln	Tyr	Glu	Glu	Asn				
									755						765				
Met	Gly	His	Pro	Gly	Ser	Arg	Lys	Phe	Asn	Ile	Thr	Glu	Arg	Val	Leu				
									770						780				
Gln	Gly	Leu	Leu	Lys	Pro	Leu	Phe	Arg	Asn	Ser	Ser	Leu	Glu	Tyr	Leu				
									785						800				

Tyr Ser Gly Cys Arg Leu Ala Ser Leu Arg Pro Glu Lys Asp Ser Ser
 805 810 815
 Ala Met Ala Val Asp Ala Ile Cys Thr His Arg Pro Asp Pro Glu Asp
 820 825 830
 Leu Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Asn Leu Thr
 835 840 845
 Asn Gly Ile Gln Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu
 850 855 860
 Tyr Val Asn Gly Phe Thr His Arg Ser Ser Met Pro Thr Thr Ser Thr
 865 870 875 880
 Pro Gly Thr Ser Thr Val Asp Val Gly Thr Ser Gly Thr Pro Ser Ser
 885 890 895
 Ser Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Met Pro Phe Thr Leu
 900 905 910
 Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr
 915 920 925
 Gly Ser Arg Lys Phe Asn Thr Met Glu Ser Val Leu Gln Gly Leu Leu
 930 935 940
 Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys
 945 950 955 960
 Arg Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala Ala Thr Gly Val
 965 970 975
 Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser Pro Gly Leu Asn
 980 985 990
 Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr Asn Asp Ile Glu
 995 1000 1005
 Glu Val Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn
 1010 1015 1020
 Gly Phe Thr His Arg Ser Phe Val Ala Pro Thr Ser Thr Leu Gly
 1025 1030 1035
 Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Ser Leu
 1040 1045 1050
 Pro Ser Pro Thr Thr Gly Val Pro Leu Leu Ile Pro Phe Thr Leu
 1055 1060 1065
 Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn Met Gly His
 1070 1075 1080
 Pro Gly Ser Arg Lys Phe Asn Ile Met Glu Arg Val Leu Gln Gly
 1085 1090 1095
 Leu Leu Ser Pro Ile Phe Lys Asn Ser Ser Val Gly Ser Leu Tyr
 1100 1105 1110

Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala
 1115 1120 1125
 Ala Thr Arg Val Asp Ala Val Cys Thr His Arg Pro Asp Pro Lys
 1130 1135 1140
 Ser Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Lys Leu Ser Gln
 1145 1150 1155
 Leu Thr His Gly Ile Ile Glu Leu Gly Pro Tyr Thr Leu Asp Arg
 1160 1165 1170
 His Ser Phe Tyr Val Asn Gly Phe Thr His Gln Ser Ser Met Thr
 1175 1180 1185
 Thr Thr Arg Thr Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser
 1190 1195 1200
 Arg Thr Pro Ala Ser Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu
 1205 1210 1215
 Leu Val Leu Phe Thr Ile Asn Phe Thr Ile Thr Asn Gln Arg Tyr
 1220 1225 1230
 Glu Glu Asn Met His His Pro Gly Ser Arg Lys Phe Asn Thr Thr
 1235 1240 1245
 Glu Arg Val Leu Gln Gly Leu Leu Arg Pro Val Phe Lys Asn Thr
 1250 1255 1260
 Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg
 1265 1270 1275
 Pro Lys Lys Asp Gly Ala Ala Thr Lys Val Asp Ala Ile Cys Thr
 1280 1285 1290
 Tyr Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu
 1295 1300 1305
 Tyr Trp Glu Leu Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly
 1310 1315 1320
 Pro Tyr Thr Gln Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr
 1325 1330 1335
 His Arg Ser Ser Val Pro Thr Thr Ser Ile Pro Gly Thr Ser Ala
 1340 1345 1350
 Val His Leu Glu Thr Ser Gly Thr Pro Ala Ser Leu Pro
 1355 1360 1365

<210> 48

<211> 1148

<212> PRT

<213> Homo sapiens

<400> 48

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Met Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu Tyr Ser Gly Cys
 1              5              10              15

Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala Ala Thr Arg Val
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Arg Glu Arg Leu Tyr Trp Lys Leu Ser Gln Leu Thr His Gly Ile Ile
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Glu Leu Gly Pro Tyr Thr Leu Asp Arg His Ser Phe Tyr Val Asn Gly
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Phe Thr His Gln Ser Ser Met Thr Thr Thr Arg Thr Pro Asp Thr Ser
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Thr Met His Leu Ala Thr Ser Arg Thr Pro Ala Ser Leu Ser Gly Pro
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Thr Thr Ala Ser Pro Leu Leu Val Leu Phe Thr Ile Asn Phe Thr Ile
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Thr Asn Gln Arg Tyr Glu Glu Asn Met His His Pro Gly Ser Arg Lys
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Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Arg Pro Val Phe
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Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu
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Leu Arg Pro Lys Lys Asp Gly Ala Ala Thr Lys Val Asp Ala Ile Cys
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Thr Tyr Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu
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Tyr Trp Glu Leu Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro
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Tyr Thr Gln Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg
225              230              235              240

Ser Ser Val Pro Thr Thr Ser Ile Pro Gly Thr Ser Ala Val His Leu
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Glu Thr Ser Gly Thr Pro Ala Ser Leu Pro Gly Pro Ser Ala Ala Ser
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Pro Leu Leu Val Leu Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Arg
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 Ser Gly Ser Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Glu Ala
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 Thr Gly Val Asp Ala Ile Cys Thr His Arg Pro Asp Pro Thr Gly Pro
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 Gly Leu Asp Arg Glu Gln Leu Tyr Leu Glu Leu Ser Gln Leu Thr His
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 Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr
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 Val Val Ser Glu Glu Pro Phe Thr Leu Asn Phe Thr Ile Asn Asn Leu
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 Arg Tyr Met Ala Asp Met Gly Gln Pro Gly Ser Leu Lys Phe Asn Ile
 580 585 590
 Thr Asp Asn Val Met Lys His Leu Leu Ser Pro Leu Phe Gln Arg Ser

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 Val Thr Glu Met Glu Ser Ser Val Tyr Gln Pro Thr Ser Ser Ser Ser
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Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 645 650 655		
Lys Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu 660 665 670		
Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu 675 680 685		
Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp 690 695 700		
Arg Gly Ser Leu Tyr Val Asn Gly Phe Thr His Arg Asn Phe Val Pro 705 710 715 720		
Ile Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Gly Thr Ser Glu 725 730 735		
Thr Pro Ser Ser Leu Pro Arg Pro Ile Val Pro Gly Pro Leu Leu Ile 740 745 750		
Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn 755 760 765		
Met Gly His Pro Gly Ser Arg Lys Phe Asn Ile Thr Glu Arg Val Leu 770 775 780		
Gln Gly Leu Leu Lys Pro Leu Phe Arg Asn Ser Ser Leu Glu Tyr Leu 785 790 795 800		
Tyr Ser Gly Cys Arg Leu Ala Ser Leu Arg Pro Glu Lys Asp Ser Ser 805 810 815		
Ala Met Ala Val Asp Ala Ile Cys Thr His Arg Pro Asp Pro Glu Asp 820 825 830		
Leu Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Asn Leu Thr 835 840 845		
Asn Gly Ile Gln Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu 850 855 860		
Tyr Val Asn Gly Phe Thr His Arg Ser Ser Met Pro Thr Thr Ser Thr 865 870 875 880		
Pro Gly Thr Ser Thr Val Asp Val Gly Thr Ser Gly Thr Pro Ser Ser 885 890 895		
Ser Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Met Pro Phe Thr Leu 900 905 910		

Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr
 915 920 925
 Gly Ser Arg Lys Phe Asn Thr Met Glu Ser Val Leu Gln Gly Leu Leu
 930 935 940
 Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys
 945 950 955 960
 Arg Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala Ala Thr Gly Val
 965 970 975
 Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser Pro Gly Leu Asn
 980 985 990
 Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr Asn Asp Ile Glu
 995 1000 1005
 Glu Val Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn
 1010 1015 1020
 Gly Phe Thr His Arg Ser Phe Val Ala Pro Thr Ser Thr Leu Gly
 1025 1030 1035
 Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Ser Leu
 1040 1045 1050
 Pro Ser Pro Thr Thr Gly Val Pro Leu Leu Ile Pro Phe Thr Leu
 1055 1060 1065
 Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn Met Gly His
 1070 1075 1080
 Pro Gly Ser Arg Lys Phe Asn Ile Met Glu Arg Val Leu Gln Gly
 1085 1090 1095
 Leu Leu Ser Pro Ile Phe Lys Asn Ser Ser Val Gly Ser Leu Tyr
 1100 1105 1110
 Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala
 1115 1120 1125
 Ala Thr Arg Val Asp Ala Val Cys Thr His Arg Pro Asp Pro Lys
 1130 1135 1140
 Ser Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Lys Leu Ser Gln
 1145 1150 1155
 Leu Thr His Gly Ile Ile Glu Leu Gly Pro Tyr Thr Leu Asp Arg
 1160 1165 1170
 His Ser Phe Tyr Val Asn Gly Phe Thr His Gln Ser Ser Met Thr
 1175 1180 1185
 Thr Thr Arg Thr Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser
 1190 1195 1200
 Arg Thr Pro Ala Ser Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu
 1205 1210 1215

Leu Val 1220	Leu Phe Thr Ile 1225	Asn Phe Thr Ile Thr 1230	Gln Arg Tyr
Glu Glu 1235	Asn Met His His 1240	Gly Ser Arg Lys Phe 1245	Asn Thr Thr
Glu Arg 1250	Val Leu Gln Gly Leu 1255	Leu Arg Pro Val Phe 1260	Lys Asn Thr
Ser Val 1265	Gly Pro Leu Tyr Ser 1270	Gly Cys Arg Leu Thr 1275	Leu Leu Arg
Pro Lys 1280	Lys Asp Gly Ala Ala 1285	Thr Lys Val Asp Ala 1290	Ile Cys Thr
Tyr Arg 1295	Pro Asp Pro Lys Ser 1300	Pro Gly Leu Asp Arg 1305	Glu Gln Leu
Tyr Trp 1310	Glu Leu Ser Gln Leu 1315	Thr His Ser Ile Thr 1320	Glu Leu Gly
Pro Tyr 1325	Thr Gln Asp Arg Asp 1330	Ser Leu Tyr Val Asn 1335	Gly Phe Thr
His Arg 1340	Ser Ser Val Pro Thr 1345	Thr Ser Ile Pro Gly 1350	Thr Ser Ala
Val His 1355	Leu Glu Thr Ser Gly 1360	Thr Pro Ala Ser Leu 1365	Pro Gly Pro
Ser Ala 1370	Ala Ser Pro Leu Leu 1375	Val Leu Phe Thr Leu 1380	Asn Phe Thr
Ile Thr 1385	Asn Leu Arg Tyr Glu 1390	Glu Asn Met Gln His 1395	Pro Gly Ser
Arg Lys 1400	Phe Asn Thr Thr Glu 1405	Arg Val Leu Gln Gly 1410	Leu Leu Arg
Ser Leu 1415	Phe Lys Ser Thr Ser 1420	Val Gly Pro Leu Tyr 1425	Ser Gly Cys
Arg Leu 1430	Thr Leu Leu Arg Pro 1435	Glu Lys Asp Gly Thr 1440	Ala Thr Gly
Val Asp 1445	Ala Ile Cys Thr His 1450	His Pro Asp Pro Lys 1455	Ser Pro Arg
Leu Asp 1460	Arg Glu Gln Leu Tyr 1465	Trp Glu Leu Ser Gln 1470	Leu Thr His
Asn Ile 1475	Thr Glu Leu Gly His 1480	Tyr Ala Leu Asp Asn 1485	Asp Ser Leu
Phe Val 1490	Asn Gly Phe Thr His 1495	Arg Ser Ser Val Ser 1500	Thr Thr Ser
Thr Pro	Gly Thr Pro Thr Val	Tyr Leu Gly Ala Ser	Lys Thr Pro

1505	1510	1515
Ala Ser Ile Phe Gly Pro Ser	Ala Ala Ser His Leu	Leu Ile Leu
1520	1525	1530
Phe Thr Leu Asn Phe Thr Ile	Thr Asn Leu Arg Tyr	Glu Glu Asn
1535	1540	1545
Met Trp Pro Gly Ser Arg Lys	Phe Asn Thr Thr Glu	Arg Val Leu
1550	1555	1560
Gln Gly Leu Leu Arg Pro Leu	Phe Lys Asn Thr Ser	Val Gly Pro
1565	1570	1575
Leu Tyr Ser Gly Ser Arg Leu	Thr Leu Leu Arg Pro	Glu Lys Asp
1580	1585	1590
Gly Glu Ala Thr Gly Val Asp	Ala Ile Cys Thr His	Arg Pro Asp
1595	1600	1605
Pro Thr Gly Pro Gly Leu Asp	Arg Glu Gln Leu Tyr	Leu Glu Leu
1610	1615	1620
Ser Gln Leu Thr His Ser Ile	Thr Glu Leu Gly Pro	Tyr Thr Leu
1625	1630	1635
Asp Arg Asp Ser Leu Tyr Val	Asn Gly Phe Thr His	Arg Ser Ser
1640	1645	1650
Val Pro Thr Thr Ser Thr Gly	Val Val Ser Glu Glu	Pro Phe Thr
1655	1660	1665
Leu Asn Phe Thr Ile Asn Asn	Leu Arg Tyr Met Ala	Asp Met Gly
1670	1675	1680
Gln Pro Gly Ser Leu Lys Phe	Asn Ile Thr Asp Asn	Val Met Lys
1685	1690	1695
His Leu Leu Ser Pro Leu Phe	Gln Arg Ser Ser Leu	Gly Ala Arg
1700	1705	1710
Tyr Thr Gly Cys Arg Val Ile	Ala Leu Arg Ser Val	Lys Asn Gly
1715	1720	1725
Ala Glu Thr Arg Val Asp Leu	Leu Cys Thr Tyr Leu	Gln Pro Leu
1730	1735	1740
Ser Gly Pro Gly Leu Pro Ile	Lys Gln Val Phe His	Glu Leu Ser
1745	1750	1755
Gln Gln Thr His Gly Ile Thr	Arg Leu Gly Pro Tyr	Ser Leu Asp
1760	1765	1770
Lys Asp Ser Leu Tyr Leu Asn	Gly Tyr Asn Glu Pro	Gly Leu Asp
1775	1780	1785
Glu Pro Pro Thr Thr Pro Lys	Pro Ala Thr Thr Phe	Leu Pro Pro
1790	1795	1800

Leu Ser 1805	Glu Ala Thr Thr	Ala 1810	Met Gly Tyr His	Leu 1815	Lys Thr Leu
Thr Leu 1820	Asn Phe Thr Ile	Ser 1825	Asn Leu Gln Tyr	Ser 1830	Pro Asp Met
Gly Lys 1835	Gly Ser Ala Thr	Phe 1840	Asn Ser Thr Glu	Gly 1845	Val Leu Gln
His Leu 1850	Leu Arg Pro Leu	Phe 1855	Gln Lys Ser Ser	Met 1860	Gly Pro Phe
Tyr Leu 1865	Gly Cys Gln Leu	Ile 1870	Ser Leu Arg Pro	Glu 1875	Lys Asp Gly
Ala Ala 1880	Thr Gly Val Asp	Thr 1885	Thr Cys Thr Tyr	His 1890	Pro Asp Pro
Val Gly 1895	Pro Gly Leu Asp	Ile 1900	Gln Gln Leu Tyr	Trp 1905	Glu Leu Ser
Gln Leu 1910	Thr His Gly Val	Thr 1915	Gln Leu Gly Phe	Tyr 1920	Val Leu Asp
Arg Asp 1925	Ser Leu Phe Ile	Asn 1930	Gly Tyr Ala Pro	Gln 1935	Asn Leu Ser
Ile Arg 1940	Gly Glu Tyr Gln	Ile 1945	Asn Phe His Ile	Val 1950	Asn Trp Asn
Leu Ser 1955	Asn Pro Asp Pro	Thr 1960	Ser Ser Glu Tyr	Ile 1965	Thr Leu Leu
Arg Asp 1970	Ile Gln Asp Lys	Val 1975	Thr Thr Leu Tyr	Lys 1980	Gly Ser Gln
Leu His 1985	Asp Thr Phe Arg	Phe 1990	Cys Leu Val Thr	Asn 1995	Leu Thr Met
Asp Ser 2000	Val Leu Val Thr	Val 2005	Lys Ala Leu Phe	Ser 2010	Ser Asn Leu
Asp Pro 2015	Ser Leu Val Glu	Gln 2020	Val Phe Leu Asp	Lys 2025	Thr Leu Asn
Ala Ser 2030	Phe His Trp Leu	Gly 2035	Ser Thr Tyr Gln	Leu 2040	Val Asp Ile
His Val 2045	Thr Glu Met Glu	Ser 2050	Ser Val Tyr Gln	Pro 2055	Thr Ser Ser
Ser Ser 2060	Thr Gln His Phe	Tyr 2065	Leu Asn Phe Thr	Ile 2070	Thr Asn Leu
Pro Tyr 2075	Ser Gln Asp Lys	Ala 2080	Gln Pro Gly Thr	Thr 2085	Asn Tyr Gln
Arg Asn 2090	Lys Arg Asn Ile	Glu 2095	Asp Ala Leu Asn	Gln 2100	Leu Phe Arg

Asn	Ser	Ser	Ile	Lys	Ser	Tyr	Phe	Ser	Asp	Cys	Gln	Val	Ser	Thr
2105						2110					2115			
Phe	Arg	Ser	Val	Pro	Asn	Arg	His	His	Thr	Gly	Val	Asp	Ser	Leu
2120						2125					2130			
Cys	Asn	Phe	Ser	Pro	Leu	Ala	Arg	Arg	Val	Asp	Arg	Val	Ala	Ile
2135						2140					2145			
Tyr	Glu	Glu	Phe	Leu	Arg	Met	Thr	Arg	Asn	Gly	Thr	Gln	Leu	Gln
2150						2155					2160			
Asn	Phe	Thr	Leu	Asp	Arg	Ser	Ser	Val	Leu	Val	Asp	Gly	Tyr	Ser
2165						2170					2175			
Pro	Asn	Arg	Asn	Glu	Pro	Leu	Thr	Gly	Asn	Ser	Asp	Leu	Pro	Phe
2180						2185					2190			
Trp	Ala	Val	Ile	Leu	Ile	Gly	Leu	Ala	Gly	Leu	Leu	Gly	Leu	Ile
2195						2200					2205			
Thr	Cys	Leu	Ile	Cys	Gly	Val	Leu	Val	Thr	Thr	Arg	Arg	Arg	Lys
2210						2215					2220			
Lys	Glu	Gly	Glu	Tyr	Asn	Val	Gln	Gln	Gln	Cys	Pro	Gly	Tyr	Tyr
2225						2230					2235			
Gln	Ser	His	Leu	Asp	Leu	Glu	Asp	Leu	Gln					
2240						2245								

<210> 51

<211> 24

<212> DNA

<213> Artificial

<220>

<223> Synthetic Primer

<400> 51

cagcagagac cagcagcagct actc

24

<210> 52

<211> 20

<212> DNA

<213> Artificial

<220>
 <223> Synthetic Primer
 <400> 52
 tccactgccca tggctgagct 20

<210> 53
 <211> 22
 <212> DNA
 <213> Artificial

<220>
 <223> Synthetic Primer
 <400> 53
 ccagcacagc tcttcccagc ac 22

<210> 54
 <211> 22
 <212> DNA
 <213> Artificial

<220>
 <223> Synthetic Primer
 <400> 54
 ggaatggctg agctgacgtc tg 22

<210> 55
 <211> 21
 <212> DNA
 <213> Artificial

<220>
 <223> Synthetic Primer
 <400> 55

cttcccagga caacctcaag g

21

<210> 56

<211> 21

<212> DNA

<213> Artificial

<220>

<223> Synthetic Primer

<400> 56

gcaggatgag tgagccacgt g

21

<210> 57

<211> 22

<212> DNA

<213> Artificial

<220>

<223> Synthetic Primer

<400> 57

gtcagatctg gtgacctcac tg

22

<210> 58

<211> 21

<212> DNA

<213> Artificial

<220>

<223> Synthetic Primer

<400> 58

gagggcactgg aaagcccaga g

21

<210> 59

<211> 25

<212> DNA

<213> Artificial

<220>

<223> Synthetic Primer

<400> 59

ctgatggcat tatggaacac atcac

25

<210> 60

<211> 22

<212> DNA

<213> Artificial

<220>

<223> Synthetic Primer

<400> 60

cccagaacga gagaccagtg ag

22

<210> 61

<211> 24

<212> DNA

<213> Artificial

<220>

<223> Synthetic Primer

<400> 61

gctgatggcg atgaatgaac actg

24

<210> 62

<211> 22

<212> DNA

<213> Artificial

<220>

<223> Synthetic Primer

<400> 62
cccagaacga gagaccagtg ag

22

<210> 63

<211> 35

<212> DNA

<213> Artificial

<220>

<223> Synthetic Primer

<400> 63
cgcggaatccg aacactgcgt ttgctggctt tgatg

35

<210> 64

<211> 23

<212> DNA

<213> Artificial

<220>

<223> Synthetic Primer

<400> 64
cctctgtgtg ctgcttcatt ggg

23

<210> 65

<211> 32

<212> DNA

<213> Artificial

<220>

<223> Synthetic Primer

<400> 65

accggatcca tggggccacac agagcctggc cc

32

<210> 66

<211> 29

<212> DNA

<213> Artificial

<220>

<223> Synthetic Primer

<400> 66

tgtaagctta ggcagggagg atggagtcc

29

<210> 67

<211> 507

<212> DNA

<213> Homo sapien

<400> 67

atgagaggat cgcacacca tcacccatcac ggatccatgg gccacacaga gcctggccct

60

ctcttgatgc cattcacttt caactttacc atcaccaacc tgcattatga ggaaacatg

120

caacaccctg gttccaggaa gttcaacacc acggagaggg ttctgcaggg tctgctcaag

180

ccctgtgtca agaacaccag tggtggccct ctgtactctg gctgcagact gaccttgctc

240

agacctgaga agcatgaggc agccactgga gtggacacca tctgtacca cgcggttgat

300

cccacggac ctggactgga cagagagcgg ctatactggg agctgagcca gctgaccaac

360

agcatcacag agctgggacc ctacaccctg gacagggaca gtctctatgt caatggcttc

420

aaccctcgga gctctgtgcc aaccaccagc actcctggga cctccacagt gcacctggca

480

acctctggga ctccatcctc cctgcct

507

<210> 68

<211> 169

<212> PRT

<213> Homo sapiens

<400> 68

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Met Arg Gly Ser His His His His His His Gly Ser Met Gly His Thr
1          5          10
Glu Pro Gly Pro Leu Leu Ile Pro Phe Thr Phe Asn Phe Thr Ile Thr
          20          25          30
Asn Leu His Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe
          35          40          45
Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys
          50          55          60
Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu
          65          70          75          80
Arg Pro Glu Lys His Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr
          85          90          95
His Arg Val Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr
          100          105          110
Trp Glu Leu Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr
          115          120          125
Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Asn Pro Arg Ser
          130          135          140
Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala
          145          150          155          160
Thr Ser Gly Thr Pro Ser Ser Leu Pro
          165

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<210> 69

<211> 909

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (1)..(909)

<223> Any "X" = any amino acid

<400> 69

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Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Met Phe Lys Asn Thr Ser
1           5           10           15

Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys
20           25           30

Lys Asp Gly Ala Ala Thr Lys Val Asp Ala Ile Cys Thr Tyr Arg Pro
35           40           45

Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu
50           55           60

Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp
65           70           75

Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr Gln Arg Ser Ser Val Pro
85           90           95

Thr Thr Ser Ile Pro Gly Thr Pro Thr Val Asp Leu Gly Thr Ser Gly
100          105          110

Thr Pro Val Ser Lys Pro Gly Pro Ser Ala Ala Ser Pro Leu Leu Ile
115          120          125

Pro Phe Thr Ile Asn Phe Thr Ile Thr Asn Leu Arg Tyr Glu Glu Asn
130          135          140

Met Gly His Pro Gly Ser Arg Lys Phe Asn Ile Met Glu Arg Val Leu
145          150          155          160

Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu
165          170          175

Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala
180          185          190

Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser
195          200          205

Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr
210          215          220

Asn Asp Ile Glu Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu
225          230          235          240

Tyr Val Asn Gly Phe Thr His Gln Ser Ser Val Ser Thr Thr Ser Thr
245          250          255

Pro Gly Thr Ser Thr Val Asp Leu Arg Thr Ser Gly Thr Pro Ser Ser
260          265          270

Leu Ser Ser Pro Thr Ile Met Ala Ala Gly Pro Leu Leu Ile Pro Phe
275          280          285

```


Thr Ile Asn Phe Thr Ile Thr Asn Leu Arg Tyr Glu Glu Asn Met His
 290 295 300
 His Pro Gly Ser Arg Lys Phe Asn Thr Met Glu Arg Val Leu Gln Gly
 305 310 315 320
 Leu Leu Met Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu Tyr Ser
 325 330 335
 Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala Ala Thr
 340 345 350
 Arg Val Asp Ala Val Cys Thr His Arg Pro Asp Pro Lys Ser Pro Gly
 355 360 365
 Leu Asp Arg Glu Arg Leu Tyr Trp Lys Leu Ser Gln Leu Thr His Gly
 370 375 380
 Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val
 385 390 395 400
 Asn Gly Phe Thr His Arg Ser Ser Met Pro Thr Thr Ser Thr Pro Gly
 405 410 415
 Thr Ser Thr Val Asp Val Gly Thr Ser Gly Thr Pro Ser Ser Ser Pro
 420 425 430
 Ser Pro Thr Thr Ala Gly Pro Leu Leu Met Pro Phe Thr Leu Asn Phe
 435 440 445
 Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr Gly Ser
 450 455 460
 Arg Lys Phe Asn Thr Met Glu Arg Val Leu Gln Gly Leu Leu Lys Pro
 465 470 475 480
 Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu
 485 490 495
 Thr Leu Leu Arg Pro Glu Lys His Gly Ala Ala Thr Gly Val Asp Ala
 500 505 510
 Ile Cys Thr Leu Arg Leu Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu
 515 520 525
 Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Ser Val Thr Glu Leu
 530 535 540
 Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr
 545 550 555 560
 His Arg Ser Ser Val Pro Thr Thr Ser Ile Pro Gly Thr Ser Ala Val
 565 570 575
 His Leu Glu Thr Ser Gly Thr Pro Ala Ser Leu Pro Gly His Thr Ala
 580 585 590
 Pro Gly Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn

595	600	605
Leu His Tyr Glu Glu Asn Met 610	Gln His Pro Gly Ser Arg Lys Phe Asn 615	
Thr Met Glu Arg Val 625	Leu Gln Gly Cys Leu Val 630	Pro Cys Ser Arg Asn 640
Thr Asn Val Gly 645	Leu Leu Tyr Ser Gly 650	Cys Arg Leu Thr Leu Leu Arg 655
Xaa Glu Lys 660	Xaa Xaa Ala Ala Thr Xaa Val Asp 665	Xaa Xaa Cys Xaa Xaa 670
Xaa Xaa Asp 675	Pro Xaa Xaa Pro Gly Leu Asp Arg 680	Glu Xaa Leu Tyr Trp 685
Glu Leu Ser Xaa 690	Leu Thr Xaa Xaa Ile Xaa Glu 695	Leu Gly Pro Tyr Thr 700
Leu Asp Arg Asn Ser 705	Leu Tyr Val Asn Gly Phe 710	Thr His Arg Ser Ser 715 720
Val Ala Pro Thr 725	Ser Thr Pro Gly Thr 730	Ser Thr Val Asp Leu Gly Thr 735
Ser Gly Thr 740	Pro Ser Ser Leu Pro Ser 745	Pro Thr Thr Val Pro Leu Leu 750
Val Pro Phe Thr 755	Leu Asn Phe Thr Ile Thr Asn 760	Leu Gln Tyr Gly Glu 765
Asp Met Arg His 770	Pro Gly Ser Arg Lys Phe Asn 775	Thr Thr Glu Arg Val 780
Leu Gln Gly Leu Leu 785	Gly Pro Leu Phe Lys Asn 790	Ser Ser Val Gly Pro 795 800
Leu Tyr Ser Gly Cys 805	Arg Leu Ile Ser Leu Arg 810	Ser Glu Lys Asp Gly 815
Ala Ala Thr Gly Val Asp 820	Ala Ile Cys Thr His 825	His Leu Asn Pro Gln 830
Ser Pro Gly Leu Asp Arg 835	Glu Gln Leu Tyr Trp 840	Gln Leu Ser Gln Val 845
Thr Asn Gly Ile Lys 850	Glu Leu Gly Pro Tyr Thr 855	Leu Asp Arg Asn Ser 860
Leu Tyr Val Asn Gly Phe 865	Thr Thr His Arg Ser Ser 870	Gly Leu Thr Thr Ser 875 880
Thr Pro Trp Thr 885	Ser Thr Val Asp Leu Gly 890	Ser Gly Thr Pro Ser 895
Pro Val Pro Ser 900	Pro Thr Thr Ala Gly 905	Pro Leu Leu Ile

<210> 70

<211> 525

<212> PRT

<213> Homo sapiens

<400> 70

Gln Gly Leu Leu Gly Pro Met Phe Lys Asn Thr Ser Val Gly Leu Leu
1 5 10 15

Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Arg Gly Ala
20 25 30

Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu Asp Pro Leu Asn
35 40 45

Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr
50 55 60

Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp Arg Gly Ser Leu
65 70 75 80

Tyr Val Asn Gly Phe Thr His Arg Asn Phe Val Pro Ile Thr Ser Thr
85 90 95

Pro Gly Thr Ser Thr Val His Leu Gly Thr Ser Glu Thr Pro Ser Ser
100 105 110

Leu Pro Arg Pro Ile Val Pro Gly Pro Leu Leu Val Pro Phe Thr Leu
115 120 125

Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Ala Met Arg His Pro
130 135 140

Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu
145 150 155 160

Arg Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu Tyr Ser Gly Cys
165 170 175

Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala Ala Thr Arg Val
180 185 190

Asp Ala Ala Cys Thr Tyr Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp
195 200 205

Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His Ser Ile Thr
210 215 220

Glu Leu Gly Pro Tyr Thr Leu Asp Arg Val Ser Leu Tyr Val Asn Gly
225 230 235 240

Phe Asn Pro Arg Ser Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser
245 250 255

Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Ser Leu Pro Gly His
 260 265 270
 Thr Ala Pro Val Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile
 275 280 285
 Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro Gly Ser Arg Lys
 290 295 300
 Phe Asn Thr Met Glu Arg Val Leu Gln Gly Leu Leu Arg Pro Leu Phe
 305 310 315 320
 Lys Asn Thr Ser Ile Gly Pro Leu Tyr Ser Ser Cys Arg Leu Thr Leu
 325 330 335
 Leu Arg Pro Glu Lys Asp Lys Ala Ala Thr Arg Val Asp Ala Ile Cys
 340 345 350
 Thr His His Pro Asp Pro Gln Ser Pro Gly Leu Asn Arg Glu Gln Leu
 355 360 365
 Tyr Trp Glu Leu Ser Gln Leu Thr His Gly Ile Thr Glu Leu Gly Pro
 370 375 380
 Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val Asp Gly Phe Thr His Trp
 385 390 395 400
 Ser Pro Ile Pro Thr Thr Ser Thr Pro Gly Thr Ser Ile Val Asn Leu
 405 410 415
 Gly Thr Ser Gly Ile Pro Pro Ser Leu Pro Glu Thr Thr Ala Thr Gly
 420 425 430
 Pro Leu Leu Ile Pro Phe Thr Pro Asn Phe Thr Ile Thr Asn Leu Gln
 435 440 445
 Tyr Glu Glu Asp Met Arg Arg Thr Gly Ser Arg Lys Phe Asn Thr Met
 450 455 460
 Glu Arg Val Leu Gln Gly Leu Leu Ser Pro Ile Phe Lys Asn Ser Ser
 465 470 475 480
 Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu
 485 490 495
 Lys Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro
 500 505 510
 Asn Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu Tyr
 515 520 525

<210> 71

<211> 594

<212> PRT

<213> Homo sapiens

<400> 71

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Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser
1          5          10          15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu
20          25          30

Lys Asp Gly Val Ala Thr Arg Val Asp Ala Ile Cys Thr His Arg Pro
35          40          45

Asp Pro Lys Ile Pro Gly Leu Asp Arg Gln Gln Leu Tyr Trp Glu Leu
50          55          60

Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp
65          70          75

Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr Gln Arg Ser Ser Val Pro
85          90          95

Thr Thr Ser Thr Pro Gly Thr Phe Thr Val Gln Pro Glu Thr Ser Glu
100         105         110

Thr Pro Ser Ser Leu Pro Gly Pro Thr Ala Thr Gly Pro Val Leu Leu
115         120         125

Pro Phe Thr Leu Asn Phe Thr Ile Ile Asn Leu Gln Tyr Glu Glu Asp
130         135         140

Met His Arg Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu
145         150         155         160

Gln Gly Leu Leu Met Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu
165         170         175

Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Gln Glu Ala
180         185         190

Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu Asp Pro Ser Glu
195         200         205

Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr
210         215         220

Asn Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu
225         230         235         240

Tyr Val Asn Gly Phe Thr His Ser Gly Val Leu Cys Pro Pro Pro Ser
245         250         255

Ile Leu Gly Ile Phe Thr Val Gln Pro Glu Thr Phe Glu Thr Pro Ser
260         265         270

Ser Leu Pro Gly Pro Thr Ala Thr Gly Pro Val Leu Leu Pro Phe Thr
275         280         285

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Leu Asn Phe Thr Ile Ile Asn Leu Gln Tyr Glu Glu Asp Met His Arg
 290 295 300
 Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu
 305 310 315 320
 Leu Thr Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly
 325 330 335
 Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Gln Glu Ala Ala Thr Gly
 340 345 350
 Val Asp Thr Ile Cys Thr His Arg Val Asp Pro Ile Gly Pro Gly Leu
 355 360 365
 Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Ser Ile
 370 375 380
 Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn
 385 390 395 400
 Gly Phe Asn Pro Trp Ser Ser Val Pro Thr Thr Ser Thr Pro Gly Thr
 405 410 415
 Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Ser Leu Pro Gly
 420 425 430
 His Thr Ala Pro Val Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr
 435 440 445
 Ile Thr Asn Leu His Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg
 450 455 460
 Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu
 465 470 475 480
 Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr
 485 490 495
 Leu Leu Arg Pro Glu Lys His Gly Ala Ala Thr Gly Val Asp Ala Ile
 500 505 510
 Cys Thr His Arg Leu Asp Pro Lys Ser Pro Gly Val Asp Arg Glu Gln
 515 520 525
 Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly
 530 535 540
 Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His
 545 550 555 560
 Trp Ile Pro Val Pro Thr Ser Ser Thr Pro Gly Thr Ser Thr Val Asp
 565 570 575
 Leu Gly Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr Thr Ala Gly
 580 585 590
 Pro Leu

<210> 72

<211> 424

<212> PRT

<213> Homo sapiens

<400> 72

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Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg
1          5          10          15

Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Tyr Thr His
20          25          30

Arg Leu Asp Pro Lys Ser Pro Gly Val Asp Arg Glu Gln Leu Tyr Trp
35          40          45

Glu Leu Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr
50          55          60

Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Gln Thr Ser
65          70          75          80

Ala Pro Asn Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Thr
85          90          95

Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr Ser Ala Gly Pro Leu
100          105          110

Leu Ile Pro Phe Thr Ile Asn Phe Thr Ile Thr Asn Leu Arg Tyr Glu
115          120          125

Glu Asn Met His His Pro Gly Ser Arg Lys Phe Asn Thr Met Glu Arg
130          135          140

Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly
145          150          155          160

Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp
165          170          175

Gly Val Ala Thr Arg Val Asp Ala Ile Cys Thr His Arg Pro Asp Pro
180          185          190

Lys Ile Pro Gly Leu Asp Arg Gln Gln Leu Tyr Trp Glu Leu Ser Gln
195          200          205

Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp
210          215          220

Ser Leu Tyr Val Asn Gly Phe Thr Gln Arg Ser Ser Val Pro Thr Thr
225          230          235          240

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Ser Thr Pro Gly Thr Phe Thr Val Gln Pro Glu Thr Ser Glu Thr Pro
 245 250 255
 Ser Ser Leu Pro Gly Pro Thr Ala Thr Gly Pro Val Leu Leu Pro Phe
 260 265 270
 Thr Leu Asn Phe Thr Ile Ile Asn Leu Gln Tyr Glu Glu Asp Met His
 275 280 285
 Arg Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly
 290 295 300
 Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser
 305 310 315 320
 Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys His Gly Ala Ala Thr
 325 330 335
 Gly Val Asp Ala Ile Cys Thr Leu Arg Leu Asp Pro Thr Gly Pro Gly
 340 345 350
 Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Ser
 355 360 365
 Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val
 370 375 380
 Asn Gly Phe Asn Pro Trp Ser Ser Val Pro Thr Thr Ser Thr Pro Gly
 385 390 395 400
 Thr Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Ser Leu Pro
 405 410 415
 Gly His Thr Ala Pro Val Pro Leu
 420

<210> 73

<211> 286

<212> PRT

<213> Homo sapiens

<400> 73

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser
 1 5 10 15
 Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu
 20 25 30
 Lys Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu
 35 40 45
 Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu
 50 55 60

Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp
65 70 75 80

Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Pro
85 90 95

Thr Thr Ser Ile Pro Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly
100 105 110

Thr Pro Ala Ser Leu Pro Gly His Thr Ala Pro Gly Pro Leu Leu Val
115 120 125

Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp
130 135 140

Met Arg His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu
145 150 155 160

Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu
165 170 175

Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Arg Gly Ala
180 185 190

Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu Asp Pro Leu Asn
195 200 205

Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr
210 215 220

Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp Arg Gly Ser Leu
225 230 235 240

Tyr Val Asn Gly Phe Thr His Arg Asn Phe Val Pro Ile Thr Ser Thr
245 250 255

Pro Gly Thr Ser Thr Val His Leu Gly Thr Ser Glu Thr Pro Ser Ser
260 265 270

Leu Pro Arg Pro Ile Val Pro Gly Pro Leu Leu Ile Pro Phe
275 280 285

<210> 74

<211> 286

<212> PRT

<213> Homo sapiens

<400> 74

Glu Arg Val Leu Gln Gly Leu Leu Arg Pro Val Phe Lys Asn Thr Ser
1 5 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys

20 25 30
 Lys Asp Gly Ala Ala Thr Lys Val Asp Ala Ile Cys Thr Tyr Arg Pro
 35 40 45
 Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu
 50 55 60
 Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp
 65 70 75 80
 Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr Gln Arg Ser Ser Val Pro
 85 90 95
 Thr Thr Ser Ile Pro Gly Thr Pro Thr Val Asp Leu Gly Thr Ser Gly
 100 105 110
 Thr Pro Val Ser Lys Pro Gly Pro Ser Ala Ala Ser Pro Leu Leu Val
 115 120 125
 Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp
 130 135 140
 Met His Arg Pro Gly Ser Arg Lys Phe Asn Ala Thr Glu Arg Val Leu
 145 150 155 160
 Gln Gly Leu Leu Ser Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu
 165 170 175
 Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu Lys Asp Gly Ala
 180 185 190
 Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro Asn Pro Lys Arg
 195 200 205
 Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr
 210 215 220
 His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp Arg Asp Ser Leu
 225 230 235 240
 Tyr Val Asn Gly Phe Thr His Gln Ser Ser Met Thr Thr Thr Arg Thr
 245 250 255
 Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser Arg Thr Pro Ala Ser
 260 265 270
 Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu Ile Pro Phe
 275 280 285
 <210> 75
 <211> 286
 <212> PRT
 <213> Homo sapiens

<400> 75

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser
 1 5 10 15
 Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu
 20 25 30
 Lys Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu
 35 40 45
 Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu
 50 55 60
 Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp
 65 70 75 80
 Arg Gly Ser Leu Tyr Val Asn Gly Phe Ser Arg Gln Ser Ser Met Thr
 85 90 95
 Thr Thr Arg Thr Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser Arg
 100 105 110
 Thr Pro Ala Ser Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu Ile
 115 120 125
 Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn
 130 135 140
 Met Gly His Pro Gly Ser Arg Lys Phe Asn Ile Met Glu Arg Val Leu
 145 150 155 160
 Gln Gly Leu Leu Asn Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu
 165 170 175
 Tyr Ser Gly Cys Arg Leu Thr Ser Leu Lys Pro Glu Lys Asp Gly Ala
 180 185 190
 Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro Asn Pro Lys Arg
 195 200 205
 Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr
 210 215 220
 His Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu
 225 230 235 240
 Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Ala Pro Thr Ser Thr
 245 250 255
 Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Ser
 260 265 270
 Leu Pro Ser Pro Thr Thr Ala Val Pro Leu Leu Ile Pro Phe
 275 280 285

<210> 76

<211> 286

<212> PRT

<213> Homo sapiens

<400> 76

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Arg Asn Ser Ser
1 5 10 15

Leu Glu Tyr Leu Tyr Ser Gly Cys Arg Leu Ala Ser Leu Arg Pro Glu
20 25 30

Lys Asp Ser Ser Ala Met Ala Val Asp Ala Ile Cys Thr His Arg Pro
35 40 45

Asp Pro Glu Asp Leu Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu
50 55 60

Ser Asn Leu Thr Asn Gly Ile Gln Glu Leu Gly Pro Tyr Thr Leu Asp
65 70 75 80

Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Gly Leu
85 90 95

Thr Thr Ser Thr Pro Trp Thr Ser Thr Val Asp Leu Gly Thr Ser Gly
100 105 110

Thr Pro Ser Pro Val Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Ile
115 120 125

Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn
130 135 140

Met Gly His Pro Gly Ser Arg Lys Phe Asn Ile Met Glu Arg Val Leu
145 150 155 160

Gln Gly Leu Leu Met Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu
165 170 175

Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala
180 185 190

Ala Thr Arg Val Asp Ala Val Cys Thr Gln Arg Pro Asp Pro Lys Ser
195 200 205

Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Lys Leu Ser Gln Leu Thr
210 215 220

His Gly Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg His Ser Leu
225 230 235 240

Tyr Val Asn Gly Leu Thr His Gln Ser Ser Met Thr Thr Thr Arg Thr
245 250 255

Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser Arg Thr Pro Ala Ser
260 265 270

Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu Ile Pro Phe
275 280 285

<210> 77

<211> 288

<212> PRT

<213> Homo sapiens

<400> 77

Glu Arg Val Leu Gln Gly Leu Leu Ser Pro Ile Ser Lys Asn Ser Ser
1 5 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu
20 25 30

Lys Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro
35 40 45

Asn Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu
50 55 60

Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp
65 70 75 80

Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Gln Asn Ser Val Pro
85 90 95

Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr Gly
100 105 110

Thr Pro Ser Ser Phe Pro Gly His Thr Glu Pro Gly Pro Leu Leu Ile
115 120 125

Pro Phe Thr Val Asn Phe Thr Ile Thr Asn Leu Arg Tyr Glu Glu Asn
130 135 140

Met His His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu
145 150 155 160

Gln Gly Leu Leu Arg Pro Val Phe Lys Asn Thr Ser Val Gly Pro Leu
165 170 175

Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala
180 185 190

Ala Thr Lys Val Asp Ala Ile Cys Thr Tyr Arg Pro Asp Pro Lys Ser
195 200 205

Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr

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210                215                220
Asn Asp Ile Glu Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu
225                230                235                240

Tyr Val Asn Gly Phe Thr His Gln Ser Ser Val Ser Thr Thr Ser Thr
                245                250                255

Pro Gly Thr Ser Thr Val Asp Leu Arg Thr Ser Gly Thr Pro Ser Ser
                260                265                270

Leu Ser Ser Pro Thr Ile Met Ala Ala Gly Pro Leu Leu Ile Pro Phe
275                280                285

<210> 78
<211> 597
<212> PRT
<213> Homo sapiens

<400> 78

Glu Arg Val Leu His Gly Leu Leu Thr Pro Leu Phe Lys Asn Thr Arg
1                5                10                15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu
20                25                30

Lys Gln Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Val
35                40                45

Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu
50                55                60

Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp
65                70                75                80

Arg Asp Ser Leu Tyr Val Asn Gly Phe Asn Pro Trp Ser Ser Val Pro
85                90                95

Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala Thr Ser Gly
100               105               110

Thr Pro Ser Ser Leu Pro Gly His Thr Ala Pro Val Pro Leu Leu Ile
115               120               125

Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu His Tyr Glu Glu Asn
130               135               140

Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu
145               150               155               160

Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu
165               170               175

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Tyr Ser Gly Cys Arg Leu Thr Leu Phe Lys Pro Glu Lys His Glu Ala
 180 185 190
 Ala Thr Gly Val Asp Ala Ile Cys Thr Leu Arg Leu Asp Pro Thr Gly
 195 200 205
 Pro Gly Leu Asp Arg Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn
 210 215 220
 Ser Val Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr
 225 230 235 240
 Val Asn Gly Phe Thr His Arg Ser Ser Val Pro Thr Thr Ser Ile Pro
 245 250 255
 Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly Thr Pro Ala Ser Leu
 260 265 270
 Pro Gly His Thr Ala Pro Gly Pro Leu Leu Ile Pro Phe Thr Leu Asn
 275 280 285
 Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr Gly
 290 295 300
 Ser Arg Lys Phe Asn Thr Met Glu Arg Val Leu Gln Gly Leu Leu Lys
 305 310 315 320
 Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg
 325 330 335
 Leu Thr Leu Leu Arg Pro Glu Lys Arg Gly Ala Ala Thr Gly Val Asp
 340 345 350
 Thr Ile Cys Thr His Arg Leu Asp Pro Leu Asn Pro Gly Leu Asp Arg
 355 360 365
 Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr Arg Gly Ile Ile Glu
 370 375 380
 Leu Gly Pro Tyr Leu Leu Asp Arg Gly Ser Leu Tyr Val Asn Gly Phe
 385 390 395 400
 Thr His Arg Asn Phe Val Pro Ile Thr Ser Thr Pro Gly Thr Ser Thr
 405 410 415
 Val His Leu Gly Thr Ser Glu Thr Pro Ser Ser Leu Pro Arg Pro Ile
 420 425 430
 Val Pro Gly Pro Leu Leu Ile Pro Phe Thr Ile Asn Phe Thr Ile Thr
 435 440 445
 Asn Leu Arg Tyr Glu Glu Asn Met His His Pro Gly Ser Arg Lys Phe
 450 455 460
 Asn Ile Met Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Leu Phe Lys
 465 470 475 480
 Asn Ser Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Ile Ser Leu
 485 490 495

Arg Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr
 500 505 510
 His His Leu Asn Pro Gln Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr
 515 520 525
 Trp Gln Leu Ser Gln Met Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr
 530 535 540
 Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser
 545 550 555 560
 Ser Gly Leu Thr Thr Ser Thr Pro Trp Thr Ser Thr Val Asp Leu Gly
 565 570 575
 Thr Ser Gly Thr Pro Ser Pro Val Pro Ser Pro Thr Thr Ala Gly Pro
 580 585 590
 Leu Leu Ile Pro Phe
 595
 <210> 79
 <211> 420
 <212> PRT
 <213> Homo sapiens
 <400> 79
 Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu Lys
 1 5 10 15
 Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro Asn
 20 25 30
 Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser
 35 40 45
 Gln Leu Thr His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp Arg
 50 55 60
 Asp Ser Leu Tyr Val Asn Gly Phe Thr His Gln Asn Ser Val Pro Thr
 65 70 75 80
 Thr Ser Thr Pro Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr Gly Thr
 85 90 95
 Pro Ser Ser Phe Pro Gly His Thr Glu Pro Gly Pro Leu Leu Ile Pro
 100 105 110
 Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn Met
 115 120 125
 Gly His Pro Gly Ser Arg Lys Phe Asn Ile Thr Glu Ser Val Leu Gln

130	135	140
Gly Leu Leu Thr Pro Leu Phe Lys Asn Ser Ser Val Gly Pro Leu Tyr		
145	150	155
Ser Gly Cys Arg Leu Ile Ser Leu Arg Ser Glu Lys Asp Gly Ala Ala		
	165	170
Thr Gly Val Asp Ala Ile Cys Thr His His Leu Asn Pro Gln Ser Pro		
	180	185
Gly Leu Asp Arg Glu Gln Leu Tyr Trp Gln Leu Ser Gln Met Thr Asn		
	195	200
Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr		
	210	215
Val Asn Gly Phe Thr His Arg Ser Leu Gly Leu Thr Thr Ser Thr Pro		
	225	230
Trp Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Pro Val		
	245	250
Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Ile Pro Phe Thr Leu Asn		
	260	265
Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn Met Gly His Pro Gly		
	275	280
Ser Arg Lys Phe Asn Ile Met Glu Arg Val Leu Gln Gly Leu Leu Arg		
	290	295
Pro Val Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg		
	305	310
Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala Ala Thr Lys Val Asp		
	325	330
Ala Ile Cys Thr Tyr Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg		
	340	345
Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His Ser Ile Thr Glu		
	355	360
Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe		
	370	375
Thr Gln Arg Ser Ser Val Pro Thr Thr Ser Ile Pro Gly Thr Pro Thr		
	385	390
Val Asp Leu Gly Thr Ser Gly Thr Pro Val Ser Lys Pro Gly Pro Ser		
	405	410
Ala Ala Ser Pro		
	420	

<210> 80

<211> 479

<212> PRT

<213> Homo sapiens

<400> 80

Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr Asn Asp Ile Glu Glu Leu
1 5 10 15

Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr
20 25 30

His Gln Ser Ser Val Ser Thr Thr Ser Thr Pro Gly Thr Ser Thr Val
35 40 45

Asp Leu Arg Thr Ser Gly Thr Pro Ser Ser Leu Ser Ser Pro Thr Ile
50 55 60

Met Ala Ala Gly Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile
65 70 75 80

Thr Asn Leu Gln Tyr Glu Glu Asn Met Gly His Pro Gly Ser Arg Lys
85 90 95

Phe Asn Ile Met Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Met Phe
100 105 110

Lys Asn Thr Ser Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu
115 120 125

Leu Arg Pro Glu Lys Asn Gly Ala Ala Thr Gly Met Asp Ala Ile Cys
130 135 140

Ser His Arg Leu Asp Pro Lys Ser Pro Gly Leu Asn Arg Glu Gln Leu
145 150 155 160

Tyr Trp Glu Leu Ser Gln Leu Thr His Gly Ile Lys Glu Leu Gly Pro
165 170 175

Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg
180 185 190

Ser Ser Val Ala Pro Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu
195 200 205

Gly Thr Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr Thr Ala Val
210 215 220

Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Lys
225 230 235 240

Tyr Glu Glu Asp Met His Cys Pro Gly Ser Arg Lys Phe Asn Thr Thr
245 250 255

Glu Arg Val Leu Gln Ser Leu Phe Gly Pro Met Phe Lys Asn Thr Ser

260 265 270
 Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Ser Glu
 275 280 285
 Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu
 290 295 300
 Asp Pro Lys Ser Leu Gly Val Asp Arg Glu Gln Leu Tyr Trp Glu Leu
 305 310 315 320
 Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp
 325 330 335
 Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Gln Thr Ser Ala Pro
 340 345 350
 Asn Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly
 355 360 365
 Thr Pro Ser Ser Leu Pro Ser Pro Thr Ser Ala Gly Pro Leu Leu Val
 370 375 380
 Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp
 385 390 395 400
 Met Arg Arg Thr Gly Ser Arg Lys Phe Asn Thr Met Glu Ser Val Leu
 405 410 415
 Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu
 420 425 430
 Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala
 435 440 445
 Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser
 450 455 460
 Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu
 465 470 475

<210> 81

<211> 5465

<212> DNA

<213> Homo sapiens

<400> 81
 cagagagcgt tgagctggga acagtgcacaa gtgcttatca agttccttca ctctcaacac 60
 ggttgacaag aactgatggc attatggaac acatcacaaa aatacccaat gaagcagcac 120
 acagaggtac cataagacca gtcaaaggcc ctcagacatc cacttcgcct gccagtccta 180
 aaggactaca cacaggaggg acaaaaagaa tggagaccac caccacagct ttgaagacca 240

ccaccacagc tttgaagacc acttccagag ccaccttgac caccagtgtc tatactccca 300
 ctttgggaac actgactccc ctcaatgcat caaggcaaat ggccagcaca atcctcacag 360
 aaatgatgat cacaacccca tatgttttcc ctgatgttcc agaaacgaca tctcatttgg 420
 ctaccagcct gggagcagaa accagcacag ctcttcccag gacaacccca tctgttctca 480
 atagagaate agagaccaca gcctcactgg tctctcgttc tggggcagag agaagtcogg 540
 ttattcaaac tctagatgtt tcttctagtg agccagatac aacagcttca tgggttatcc 600
 atctgcgaga gaccatocca actgtttcca agacaacccc caatttttcc cacagtgaat 660
 tagacactgt atcttccaca gccaccagtc atggggcaga cgtcagctca gccattccaa 720
 caaatatctc acctagttaa ctagatgcac tgacccactt ggtcactatt tcggggacag 780
 atactagtac aacattccca aactgacta agtcccaca tgaacagag acaagaacca 840
 catggctcac tcactctgca gagaccagct caactattcc cagaacaatc cccaattttt 900
 ctcatcatga atcagatgcc acaccttcaa tagccaccag tctgtgggca gaaaccagtt 960
 cagctattcc aattatgact gtctcacctg gtgcagaaga tctggtgacc tcacagggtca 1020
 ctagtctcgg gacagacaga aatatgacta ttccaaactt gactetttct cctggtgaac 1080
 caaagacgat agcotcatta gtcacccatc ctgaagcaca gacaagttcg gccattccaa 1140
 cttcaactat ctgcctgct gtatcacggt tggtagcttc aatggtcacc agtttggcgg 1200
 caaagacaag tacaactaat cgagctctga caaactcccc tgggtgaacca gctacaacag 1260
 ttctattggt cagcctcct gcacagacca gcccacagct tccctggaca acttccattt 1320
 tttccatag taaatcagac accacacctt caatgaccac cagtcattggg gcagaatcca 1380
 gttcagctgt tccaaactca actgtttcaa ctgaggtacc aggagtatgt acccctttgg 1440
 tcaccagttc tagggcagtg atcagtacaa ctattccaat tctgactctt tctcctggtg 1500
 aaccagagac cacaccttca atggccacca gtcattggga agaagccagt tctgtatttc 1560
 caactccaac tgtttcacct ggggtaccag gagtggtgac ctctctggtc actagtctca 1620
 gggcagtgac tagtacaact attccaatto tgacttttct tcttggtgaa ccagagacca 1680
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 <211> 1821
 <212> PRT
 <213> Homo sapiens

<400> 82

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 35 40 45
 Gly Pro Gln Thr Ser Thr Ser Pro Ala Ser Pro Lys Gly Leu His Thr
 50 55 60
 Gly Gly Thr Lys Arg Met Glu Thr Thr Thr Thr Ala Leu Lys Thr Thr
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 Thr Thr Ala Leu Lys Thr Thr Ser Arg Ala Thr Leu Thr Thr Ser Val
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 Tyr Thr Pro Thr Leu Gly Thr Leu Thr Pro Leu Asn Ala Ser Arg Gln
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 Met Ala Ser Thr Ile Leu Thr Glu Met Met Ile Thr Thr Pro Tyr Val
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 Ala Glu Thr Ser Thr Ala Leu Pro Arg Thr Thr Pro Ser Val Leu Asn
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 Arg Ser Pro Val Ile Gln Thr Leu Asp Val Ser Ser Ser Glu Pro Asp
 180 185 190
 Thr Thr Ala Ser Trp Val Ile His Pro Ala Glu Thr Ile Pro Thr Val
 195 200 205
 Ser Lys Thr Thr Pro Asn Phe Phe His Ser Glu Leu Asp Thr Val Ser
 210 215 220
 Ser Thr Ala Thr Ser His Gly Ala Asp Val Ser Ser Ala Ile Pro Thr
 225 230 235 240
 Asn Ile Ser Pro Ser Glu Leu Asp Ala Leu Thr Pro Leu Val Thr Ile

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His Glu Thr Glu Thr Arg Thr Thr Trp Leu Thr His Pro Ala Glu Thr 275 280 285		
Ser Ser Thr Ile Pro Arg Thr Ile Pro Asn Phe Ser His His Glu Ser 290 295 300		
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Ala Ile Pro Ile Met Thr Val Ser Pro Gly Ala Glu Asp Leu Val Thr 325 330 335		
Ser Gln Val Thr Ser Ser Gly Thr Asp Arg Asn Met Thr Ile Pro Thr 340 345 350		
Leu Thr Leu Ser Pro Gly Glu Pro Lys Thr Ile Ala Ser Leu Val Thr 355 360 365		
His Pro Glu Ala Gln Thr Ser Ser Ala Ile Pro Thr Ser Thr Ile Ser 370 375 380		
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Lys Thr Ser Thr Thr Asn Arg Ala Leu Thr Asn Ser Pro Gly Glu Pro 405 410 415		
Ala Thr Thr Val Ser Leu Val Thr His Pro Ala Gln Thr Ser Pro Thr 420 425 430		
Val Pro Trp Thr Thr Ser Ile Phe Phe His Ser Lys Ser Asp Thr Thr 435 440 445		
Pro Ser Met Thr Thr Ser His Gly Ala Glu Ser Ser Ser Ala Val Pro 450 455 460		
Thr Pro Thr Val Ser Thr Glu Val Pro Gly Val Val Thr Pro Leu Val 465 470 475 480		
Thr Ser Ser Arg Ala Val Ile Ser Thr Thr Ile Pro Ile Leu Thr Leu 485 490 495		
Ser Pro Gly Glu Pro Glu Thr Thr Pro Ser Met Ala Thr Ser His Gly 500 505 510		
Glu Glu Ala Ser Ser Ala Ile Pro Thr Pro Thr Val Ser Pro Gly Val 515 520 525		
Pro Gly Val Val Thr Ser Leu Val Thr Ser Ser Arg Ala Val Thr Ser 530 535 540		
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Ser	Arg	Ala	Val	Thr	Ser	Thr	Thr	Leu	Pro	Thr	Leu	Thr	Leu	Ser	Pro
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Thr	Ser	Leu	Val	Thr	Ser	Ser	Ser	Gly	Val	Asn	Ser	Thr	Ser	Ile	Pro
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				675			680					685			
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Ser	Ala	Val	Leu	Thr	Val	Ser	Pro	Glu	Val	Pro	Gly	Met	Val	Thr	Ser
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Leu	Val	Thr	Ser	Ser	Arg	Ala	Val	Thr	Ser	Thr	Thr	Ile	Pro	Thr	Leu
				755			760					765			
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Val	Thr	His	Pro	Ala	Glu	Ser	Ser	Ser	Thr	Leu	Pro	Arg	Thr	Thr	Ser
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 Pro Glu Ala Glu Ser Ser Ser Ala Ile Ser Thr Thr Ile Ser Pro Gly
 900 905 910
 Ile Pro Gly Val Leu Thr Ser Leu Val Thr Ser Ser Gly Arg Asp Ile
 915 920 925
 Ser Ala Thr Phe Pro Thr Val Pro Glu Ser Pro His Glu Ser Glu Ala
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 Thr Ala Ser Trp Val Thr His Pro Ala Val Thr Ser Thr Thr Val Pro
 945 950 955 960
 Arg Thr Thr Pro Asn Tyr Ser His Ser Glu Pro Asp Thr Thr Pro Ser
 965 970 975
 Ile Ala Thr Ser Pro Gly Ala Glu Ala Thr Ser Asp Phe Pro Thr Ile
 980 985 990
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 995 1000 1005
 Ser Gly Thr Asp Thr Ser Ile Thr Ile Pro Thr Leu Thr Leu Ser
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 Ser Thr Thr Thr Phe Pro Thr Leu Thr Glu Thr Pro Tyr Glu Pro
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 Glu Thr Thr Ala Ile Gln Leu Ile His Pro Ala Glu Thr Asn Thr
 1085 1090 1095
 Met Val Pro Arg Thr Thr Pro Lys Phe Ser His Ser Lys Ser Asp
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 Thr Thr Leu Pro Val Ala Ile Thr Ser Pro Gly Pro Glu Ala Ser
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 Thr Trp Leu Thr His Pro Ala Glu Thr Ser Thr Thr Val Ser Gly

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Gln Thr Ser Thr Pro Val 1295	Ser Arg Thr Thr Ser 1300	Ser Phe Ser His 1305
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Phe Pro Ala Ser Thr Val 1385	Phe Pro Gln Val Ser 1390	Glu Thr Thr Ala 1395
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Ser Arg Val Asp Leu Ser 1430	Pro Thr Ala Ser Pro 1435	Gly Val Ser Ala 1440
Lys Thr Ala Pro Leu Ser 1445	Thr His Pro Gly Thr 1450	Glu Thr Ser Thr 1455
Met Ile Pro Thr Ser Thr 1460	Leu Ser Leu Gly Leu 1465	Leu Glu Thr Thr 1470

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1490						1495					1500				
Ser	Ile	Thr	Thr	Asp	Lys	Pro	Gln	Thr	Val	Thr	Ser	Trp	Asn	Thr	
1505						1510					1515				
Glu	Thr	Ser	Pro	Ser	Val	Thr	Ser	Val	Gly	Pro	Pro	Glu	Phe	Ser	
1520						1525					1530				
Arg	Thr	Val	Thr	Gly	Thr	Thr	Met	Thr	Leu	Ile	Pro	Ser	Glu	Met	
1535						1540					1545				
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1550						1555					1560				
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1565						1570					1575				
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1580						1585					1590				
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1595						1600					1605				
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1610						1615					1620				
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Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly
1775 1780 1785

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Ser Pro Thr
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<211> 468

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<213> Homo sapiens

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catcgccctg accctgaaga cctcggactg gacagagagc gactgtactg ggagctgagc 300
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gtcaatgggtt tcaccatcag aagetctatg cccaccacca gcaactcctgg gacctccaca 420
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<211> 474

<212> DNA

<213> Homo sapiens

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gtcaatggtt tcacccatca gagctctgtg tccaccacca gcaactcctgg gacctccaca 420
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<210> 85

<211> 468

<212> DNA

<213> Homo sapiens

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<210> 86

<211> 465

<212> DNA

<213> Homo sapiens

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<400> 86
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<210> 87

<211> 468

<212> DNA

<213> Homo sapiens

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caccgtcttg accccaaaag cctgggagt gacagggagc agctatactg ggagctgagc 300
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gtcaatgggt tcacccatca gacctctgcg cccaacacca gcactcctgg gacctccaca 420
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<210> 88

<211> 468

<212> DNA

<213> Homo sapiens

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<222> (1)..(468)

<223> All N's = any nucleotide

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<400> 88
ncnnctgncc ctctcctgnt nccnttcacc ntcaacttna ccacaccaa cctgcantan 60
gnggannaca tgcnnncccc nggntccagg aagttcaaca ccacngagng ngtnctgcag 120
```

```

ggtctgctnn nncccntntt caagaacacc agtgttgcc ctctgtactc tggtgcaga 180
ctgaccttgc tcaggctcga gaaggatgga gcagccactg gagtggatgc catctgcacc 240
caccgtcttg accccaaaag ccctggagtg gacagggagc agctatactg ggagctgagc 300
cagctgacca atggcatcaa agagctgggt ccctacacc tggaacagaaa cagtctctat 360
gtcaatgggt tcacctatca gacctctgcg cccaacacca gacctcctgg gacctccaca 420
gtggaccttg ggacctcagg gactccatcc tccctcccca gccctaca 468

```

<210> 89

<211> 468

<212> DNA

<213> Homo sapiens

```

<400> 89
tctgctggcc ctctcctggt gccattcacc ctcaacttca ccatacacia cctgcagtag 60
gaggaggaca tgcattcacc aggtctccagg aagttaaca ccacggagcg ggtcctgcag 120
ggtctgcttg gtccatggt caagaacacc agtgtcgcc ttctgtactc tggtgcaga 180
ctgaccttgc tcaggcctga gaagaatggg gcagccactg gaatggatgc catctgcacc 240
caccgtcttg accccaaaag ccctggactc aacagagagc agctgtactg ggagctgagc 300
cagctgacct atggcatcaa agagctgggc ccctacacc tggaacagaa cagtctctat 360
gtcaatgggt tcacctatcg gacctctgtg gcccccacca gacctcctgg gacctccaca 420
gtggaccttg ggacctcagg gactccatcc tccctcccca gccccaca 468

```

<210> 90

<211> 468

<212> DNA

<213> Homo sapiens

```

<400> 90
acagctgttc ctctcctggt gccgttcacc ctcaacttta ccatacacia tctgcagtag 60
gggaggagaca tgcgtcacc tggtctccagg aagttaaca ccacagagag ggtcctgcag 120
ggtctgcttg gtcccttggt caagaactcc agtgtcgcc ctctgtactc tggtgcaga 180

```


ctgatctctc tcaggtctga gaaggatggg gcagccactg gagtggatgc catctgcacc	240
caccacctta accctcaaag ccctggactg gacagggagc agctgtactg gcagctgagc	300
cagatgacca atggcatcaa agagctgggc ccctacaccc tggaccggaa cagtctctac	360
gtcaatggtt tcacccatcg gagctctggg ctcaccacca gcaactcctg gacttcocaca	420
gttgaccttg gaacctcagc gactccatcc ccggtcccca gcccacaca	468

<210> 91

<211> 468

<212> DNA

<213> Homo sapiens

<400> 91 actgctggcc ctctcctggt gccattcacc ctcaacttca ccatacaccia cctgcagtat	60
gaggaggaca tgcacgcgcc tggatctagg aagttcaaca ccacagagag ggtcctgcag	120
ggtctgctta gtccatttt caagaactcc agtggtggcc ctctgtactc tggctgcaga	180
ctgacctctc tcaggccoga gaaggatggg gcagcaactg gaatggatgc tgtctgcctc	240
taccacccta atcccaaaag acctggactg gacagagagc agctgtactg ggagctaagc	300
cagctgaccc acaacatcac tgagctgggc ccctacagcc tggacaggga cagtctctat	360
gtcaatggtt tcacccatca gaactctgtg cccaccacca gtactccttg gacctccaca	420
gtgtactggg caacctctg gactccatcc tccttcoccc gccacaca	468

<210> 92

<211> 468

<212> DNA

<213> Homo sapiens

<400> 92 gagcctggcc ctctcctgat accattcact ttcaacttta ccatacaccia cctgcattat	60
gaggaaaaca tgcaacaccc tggttccagg aagttcaaca ccacggagag ggttctgcag	120
ggtctgctca agccctgtt caagaacacc agtggtggcc ctctgtactc tggctgcaga	180
ctgacctctc tcaggccoga gaaggatggg gcagcaactg gaatggatgc tgtctgcctc	240
taccacccta atcccaaaag acctgggctg gacagagagc agctgtactg ggagctaagc	300

```

cagctgaccc acaacatcac tgagctgggc ccctacagcc tggacaggga cagtctctat 360
gtcaatgggt tcaaccca caaactctgtg cccaccacca gtactcctgg gacctccaca 420
gtgtactggg caaccactgg gactccatcc tccttccccc gccacaca 468

```

<210> 93

<211> 468

<212> DNA

<213> Homo sapiens

```

<400> 93
gagcctgggc ctctcctgat accattcact ttcaacttta ccatcaccaa cctgcattat 60
gaggaataca tgcaacaccc tggttccagg aagttcaaca ccacggagag ggttctgcag 120
ggtctgtctca agcccttggt caagaacacc agtggtggcc ctctgtactc tggctgcaga 180
ctgaccttgc tcagacctga gaagcatgag gcagccactg gagtggacac catctgtacc 240
caccgcgttg atcccatcgg acctggactg gacagggagc ggctatactg ggagctgagc 300
cagctgacca acagcattac cgaactggga ccctacaccc tggacaggga cagtctctat 360
gtcaatgggt tcaacccctg gagctctgtg ccaaccacca gcactcctgg gacctccaca 420
gtgcacctgg caacctctgg gactccatcc tccttgccctg gccacaca 468

```

<210> 94

<211> 468

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)..(468)

<223> All N's = any nucleotide

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<400> 94
gccccctgtc ctctcttgat accattcacc ctcaacttta ccatcaccaa cctgcattat 60

```

```

gaggaaaaca tgcaacaccc tgggtccagg aagttcaaca ccacggagag ggtctgcag 120
ggtctgctca agcccttgtt caagaacacc agtggtggcc ctctgtactc tggctgcaga 180
ctgaccttgc tcagacctga gaagcatgag gcagccactg gaggggacac catctgtacc 240
caccgcgttg atcccatcgg acctggactg nacagngagc ngctntactg ggagctnagc 300
canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat 360
gtcaatgggt tcacctatcn ganctctgng cccaccacca gactcctgg gacctccaca 420
gtgnaentng gnacctongg gactccatcc tccntccccc gccncaca 468

```

<210> 95

<211> 468

<212> DNA

<213> Homo sapiens

```

<400> 95
tctgctggcc ctctcctggt gccattcacc ctcaacttca ccatcaccaa cctgcagtac 60
gaggaggaca tgcattaccc aggtctccagg aagttcaaca ccacggagcg ggtcctgcag 120
ggtctgcttg gtcccatgtt caagaacacc agtgctggcc ttctgtactc tggctgcaga 180
ctgaccttgc tcaggcctga gaagaatggg gcagccactg gaatggatgc catctgcagc 240
caccgtcttg accccaaaag cctctggactc gacagagagc agctgtactg ggagctgagc 300
cagctgaccc atggcatcaa agagctgggc ccctacaccc tggacaggaa cagtctctat 360
gtcaatgggt tcacctatcg gagctctgtg gcccccacca gactcctgg gacctccaca 420
gtggaccttg ggacctcagg gactccatcc tccctcccca gccccaca 468

```

<210> 96

<211> 468

<212> DNA

<213> Homo sapiens

```

<400> 96
acagctgttc ctctcctggt gccgttcacc ctcaacttta ccatcaccaa tctgcagtat 60
ggggaggaca tgcgtacccc tggctccagg aagttcaaca ccacagagag ggtcctgcag 120
ggtctgcttg gtcccttgtt caagaactcc agtgctggcc ctctgtactc tggctgcaga 180

```

```

ctgatctctc tcaggtctga gaaggatggg gcagccactg gagtggatgc catctgcacc 240
caccacctta accctcaaag ccctggactg gacaggggagc agctgtactg gcagctgagc 300
cagatgacca atggcatcaa agagctgggc ccctacaccc tggaccggaa cagtctctac 360
gtcaatggtt tcaccatcg gagctctggg ctcaccacca gcaactcctg gacttcocaca 420
gttgaccttg gaacctcagg gactccatcc cccgtcccca gcccacaca 468

```

<210> 97

<211> 468

<212> DNA

<213> Homo sapiens

```

<400> 97
actgctggcc ctctcctggt gccattcacc ctaaacttca ccatcaccaa cctgcagtat 60
gaggaggaca tgcctcgccc tggatctagg aagtccaacg ccacagagag ggtcctgcag 120
ggtctgctta gtcccatatt caagaactcc agtggtggcc ctctgtactc tggctgcaga 180
ctgacctctc tcaggcccca gaaggatggg gcagcaactg gaatggatgc tgtctgcctc 240
taccacccta atcccaaaag ccctggactg gacagagagc agctgtactg ggagctaagc 300
cagctgacct acaacatcac tgagctgggc ccctacagcc tggacaggga cagtctctat 360
gtcaatggtt tcaccatca gagctctatg acgaccacca gaactcctga tacctccaca 420
atgcacctgg caacctcgag aactccagcc tccctgtctg gacctacg 468

```

<210> 98

<211> 474

<212> DNA

<213> Homo sapiens

```

<400> 98
accgccagcc ctctcctggt gctattcaca atcaactgca ccatcaccaa cctgcagtac 60
gaggaggaca tgcctcgcac tggctccagg aagtccaaca ccatggagag tgtcctgcag 120
ggtctgctca agcccttgtt caagaacacc agtggtggcc ctctgtactc tggctgcaga 180
ttgaccttgc tcaggcccaa gaaagatggg gcagccactg gagtggatgc catctgcacc 240

```

```

caccgccttg accccaaaag ccttggaactc aacagggagc agctgtactg ggagctaagc 300
aaactgacca atgacattga agagctgggc ccctacaccc tggacaggaa cagtctctat 360
gtcaatgggt tcacccatca gagctctgtg tcacccacca gcaactcctgg gacctccaca 420
gtgggatctca gaacctcagg gactccatcc tccctctcca gccccacaat tatg 474

```

<210> 99

<211> 468

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)..(468)

<223> All N's = any nucleotide

```

<400> 99
nncnctgncc ctctcctgnt nccnttcacc ntcaacttna ccatcaceaa cctgcantan 60
gnggannaca tgcnnncgcc nggntccagg aagttcaaca ccacngagag ggtcctacag 120
ggtctgctca gcccttggtt caagaacacc agtgtcagct ctctgtactc tggttgcaga 180
ctgaccttgc tcaggcctga gaaggatggg gcagccacca gagtggatgc tgccctgcacc 240
taccgccttg atcccaaaag ccttggaactg gacagagagc aactatactg ggagctgagc 300
cagctaacc acagcatcac tgagctggga ccctacaccc tggacagggt cagtctctat 360
gtcaatgggt tcaacctctg gagctctgtg ccaaccaeca gcaactcctgg gacctccaca 420
gtgcacctgg caacctctgg gactccatcc tccctgcctg gccacaca 468

```

<210> 100

<211> 468

<212> DNA

<213> Homo sapiens

```

<400> 100
gccctgtccc ctctcttgat accattcacc ctcaacttta ccatcaceaa cctgcattat 60

```

```

gaagaaaaca tgcaacaccc tggttccagg aagttcaaca ccacggagag ggttctgcag 120
ggtctgctca agcccttggt caagagcacc agcgttggcc ctctgtactc tggctgcaga 180
ctgaccttgc tcagacctga gaaacatggg gcagccactg gagtggacgc catctgcacc 240
ctccgccttg atcccaactgg tccctggactg gacagagagc ggctatactg ggagctgagc 300
cagctgacca acagcgttac agagctgggc ccctacaccc tggacagggg cagtctctat 360
gtcaatggct tcaccacagc gagctctgtg ccaaccacca gtattcctgg gacctctgca 420
gtgcacctgg aaacctctgg gactccagcc tccctccctg gccacaca 468

```

<210> 101

<211> 468

<212> DNA

<213> Homo sapiens

```

<400> 101
gcccctggcc ctctcctggt gccattcacc ctcaacttca ctatcaccaa cctgcagtat 60
gagggtgaca tgcgtcacc tggttccagg aagttcaaca ccacggagag agtctgcag 120
ggtctgctca agcccttggt caagagcacc agtgttg gcc ctctgtactc tggctgcaga 180
ctgaccttgc tcagacctga aaaaagtggg gcagccaccg gctggacac catctgcact 240
caccgccttg accctctaaa ccttggactg gacagagagc agctatactg ggagctgagc 300
aaactgaccc gtggcatcat cgagctgggc ccctacctcc tggacagagg cagtctctat 360
gtcaatgggt tcaccatcg gaactttgtg cccatcacca gcaactcctg gacctccaca 420
gtcacacctg gaacctctga aactccatcc tccctacctc gaccacata 468

```

<210> 102

<211> 468

<212> DNA

<213> Homo sapiens

```

<400> 102
gtgcttg gcc ctctcctggt gccattcacc ctcaacttca ccatcaccaa cttgcagtat 60
gaggaggcca tgcgacaccc tggctccagg aagttcaata ccacggagag ggtcctacag 120

```

<213> Homo sapiens

<400> 104
 actgctggcc ctctcctggg gccattcacc ctcaacttca ccatcaccaa cctgcagtat 60
 gaggaggaca tgcacgccc tggttccagg aggttcaaca ccacggagag ggttctgcag 120
 ggtctgctca cgcccttggt caagaacacc agtgttgccc ctctgtactc tggctgcaga 180
 ctgaccttgc tcagacctga gaagcaagag gcagccactg gagtggacac catctgtacc 240
 caccgcgttg atcccatcgg acctggactg gacagagagc ggctatactg ggagctgagc 300
 cagctgacca acagcatcac agagctggga ccctacacc tggataggga cagtctctat 360
 gtcaatggct tcaaccttg gagctctgtg ccaaccacca gactcctgg gacctccaca 420
 gtgcacctgg caacctctgg gactccatcc tccctgcctg gccacaca 468

<210> 105

<211> 468

<212> DNA

<213> Homo sapiens

<400> 105
 gccctgtcc ctctcttgat accattcacc ctcaacttta ccatcacaga cctgcattat 60
 gaagaaaaca tgcaacaccc tggttccagg aagttcaaca ccacggagag ggttctgcag 120
 ggtctgctca agcccttggt caagagcacc agcgttgccc ctctgtactc tggctgcaga 180
 ctgaccttgc tcagacctga gaaacatggg gcagccactg gagtggacgc catctgcacc 240
 ctccgccttg atcccaactg tcttgactg gacagagagc ggctatactg ggagctgagc 300
 cagctgacca acagcgttac agagctgggc ccctacacc tggacaggga cagtctctat 360
 gtcaatggct tcaccatcg gagctctgtg ccaaccacca gtattcctgg gacctctgca 420
 gtgcacctgg aaacctctgg gactccagcc tccctccctg gccacaca 468

<210> 106

<211> 468

<212> DNA

<213> Homo sapiens

<400> 106


```

gccccctggcc ctctcctggt gccattcacc ctcaacttca ctatcaccaa cctgcagtat      60
gaggaggaca tgcgtcacc tggttccagg aagttcagca ccacggagag agtcctgcag      120
ggtctgctca agcccttggt caagaacacc agtgtcagct ctctgtactc tggttgcaga      180
ctgaccttgc tcaggcctga gaaggatggg gcagccacca gagtggatgc tgtctgcacc      240
catcgtcctg accccaaaag ccttgactg gacagagagc ggctgtactg gaagctgagc      300
cagctgaccc acggcatcac tgagctgggc ccctacacc tggacaggca cagtctctat      360
gtcaatgggt tcacccatca gagctctatg acgaccacca gaactcctga tacctccaca      420
atgcacctgg caacctcgag aactccagcc tccctgtctg gacctacg      468

```

<210> 107

<211> 468

<212> DNA

<213> Homo sapiens

```

<400> 107
accgccagcc ctctcctggt gctattcaca attaacttca ccactactaa cctgcggtat      60
gaggagaaca tgcattcacc tggctctaga aagtttaaca ccacggagag agtccttcag      120
ggtctgctca ggcctgtggt caagaacacc agtggtggcc ctctgtactc tggctgcaga      180
ctgaccacgc tcaggcccaa gaaggatggg gcagccacca aagtggatgc catctgcacc      240
taccgccctg atcccaaaag ccttgactg gacagagagc agctatactg ggagctgagc      300
cagctaacc acagcatcac tgagctgggc ccctacacc aggacaggga cagtctctat      360
gtcaatgggt tcacccatcg gagctctgtg ccaaccacca gtattcctgg gacctctgca      420
gtgcacctgg aaacctctg gactccagcc tccctccctg gccacaca      468

```

<210> 108

<211> 468

<212> DNA

<213> Homo sapiens

```

<400> 108
gccccctggcc ctctcctggt gccattcacc ctcaacttca ctatcaccaa cctgcagtat      60
gaggaggaca tgcgtcacc tggttccagg aagttcaaca ccacggagag agtcctgcag      120

```

```

ggctctgctca agcccttggt caagagcacc agtgttggcc ctctgtactc tggtcgcaga 180
ctgaccttgc tcaggcctga aaaacgtggg gcagccaccg gcgtggacac catctgcact 240
caccgccttg accctctaaa ccaggactg gacagagagc agctatactg ggagctgagc 300
aaactgacce gtggcatcat cgagctgggc cctacctcc tggacagagg cagtctctat 360
gtcaatgggt tcaccatcg gacctctgtg cccaccacca gcaactctgg gacctcaca 420
gtggaccttg gaacctcagg gactccatc tccctcccaa gccccgca 468

```

<210> 109

<211> 465

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)..(465)

<223> All N's = any nucleotide

```

<400> 109
nennctgncc ctctctgnt ncnnttcacc ntcaacttna ccatcaccaa cctgcantan 60
gnggannaca tgcnnncocc nggntccagg aagttaaca ccacngagag ggtcctgcag 120
actctgcttg gtcctatggt caagaacacc agtgttggcc ttctgtactc tggtcgcaga 180
ctgaccttgc tcaggtcoga gaaggatgga gcagccactg gagtggatgc catctgcacc 240
caccgtcttg accccaaaag ccttgaggatg gacagggagc aactatactg ggagctgagc 300
cagctgacca atggcattaa agaactgggc cctacacccc tggacaggaa cagtctctat 360
gtcaatgggt tcaccattg gatccctgtg cccaccagca gcaactctgg gacctcaca 420
gtggaccttg ggtcaggggac tccatctctc cccccagcc ccaca 465

```

<210> 110

<211> 468

<212> DNA

<213> Homo sapiens

```

<400> 110
actgctggcc ctctcctggt gccgttcacc ctcaacttca ccatacacia cctgaagtac      60
gaggaggaca tgcattgcc ttgctccagg aagttaaca ccacagagag agtctgtcac      120
agtctgcttg gtcccatgtt caagaacacc agtggtggcc ctctgtactc tggctgcaga      180
ctgaccttgc tcaggtcga gaaggatgga gcagccactg gagggatgc catctgcacc      240
caccgtcttg acccaaaaag ccttgaggag gacaggagag agctatactg ggagctgagc      300
cagctgacca atggcatcaa agagctgggt cctacacacc tggacagaaa cagtctctat      360
gtcaatgggt tcacccatca gacctctgag ccaacacaca gacctctgag gacctccaca      420
gtggaccttg ggacctcagg gactccatcc tccctcccca gccctaca      468

```

<210> 111

<211> 465

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)..(465)

<223> All N's = any nucleotide

```

<400> 111
nennctgncc ctctcctgnt nccnttcacc ntcaacttna ccatacacia cctgcantan      60
gnggannaca tgcnncccc nggntccagg aagttaaca ccacngagng ngtnctgcag      120
ggtctgtcnn nccccntntt caagaacncc agtgtnggcc ntctgtactc tggctgcaga      180
ctgacctnnc tcaggncnga gaagnatggn gcagccactg gantggatgc catctgcanc      240
caccnncntn ancccaaaaag nccctggaactg nacagngagc ngctntactg ggagctnagc      300
cancctgacca annncatcnn ngagctgggn cctacacacc tggacaggna cagtctctat      360
gtcaatgggt tcacccattg gatccctgtg cccaccagca gacctctgag gacctccaca      420
gtggaccttg ggtcaggagc tccatcctcc ctccccagcc ccaca      465

```

<210> 112
 <211> 468
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)..(468)
 <223> All N's = any nucleotide

<400> 112
 actgctggcc ctctctctggt gccgttcacc ctcaacttca ccatcaccaa cctgaagtac 60
 gaggaggaca tgcattgcc tggctccagg aagtccaaca ccacagagag agtctcgag 120
 agtctgcttg gtcccatggt caagaacacc agtgttgccc ctctgtactc tggctgcaga 180
 ctgacctcgc tcagggtccga gaaggatgga gcagccactg gagtggatgc catctgcacc 240
 caccgtgttg accccaaaag ccctggagtg gacagggagc agctatactg ggagctgagc 300
 cagctgacca atggcatcaa agagctgggt ccctacaccc tggacagaaa cagtctctat 360
 gtcaatggtt tcacccatca gacctctgcy cccaacacca gcactcctgg gacctccaca 420
 gtgnactntg gnacctcngg gactccatcc tcctccccc gccncaca 468

<210> 113
 <211> 468
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)..(468)
 <223> All N's = any nucleotide

```

<400> 113
tctgctggcc ctctcctggt gccattcacc ctcaacttca ccatcaccaa cctgcagtac      60
gaggaggaca tgcattcacc aggtccagg aagtccaaca ccacggagcg ggtcctgcag      120
ggtctgcttg gtcccatggt caagaacacc agtgtcggcc ttctgtactc tggctgcaga      180
ctgaccttgc tcaggcctga gaagaatggg gcaaccactg gaatggatgc catctgcacc      240
caccgtcttg accccaaaag ccttggaactg nacagnagag ngetntactg ggagctnagc      300
canctgacca annncatcnn ngagctgggn cctacacccc tggacaggna cagtctctat      360
gtcaatgggt tcacccatcn gantctgng cccaccacca gactcctgg gacctccaca      420
gtgnaentng gnacctcngg gactccatcc tcctccccc gccncaca      468

```

```
<210> 114
```

```
<211> 468
```

```
<212> DNA
```

```
<213> Homo sapiens
```

```
<220>
```

```
<221> misc_feature
```

```
<222> (1)..(468)
```

```
<223> All N's = any nucleotide
```

```

<400> 114
ncnnetgncc ctctcctgnt nccttcacc ntcaacttna ccatcaccaa cctgcantan      60
gnggannaca tgennncccc nggntccagg aagtccaaca ccacngagag ggtctgcag      120
ggtctgctca aaccttggtt caggaatagc agtctggaat acctctatc aggtcgcaga      180
ctagcctcac tcaggccaga gaaggatagc tcagccatgg cagtggatgc catctgcaca      240
catcgccctg accctgaaga cctcggactg gacagagagc gactgtactg ggagctgagc      300
aatctgacaa atggcatcca ggagctgggc cctacacccc tggaccggaa cagtctctat      360
gtcaatgggt tcacccatcg aagctctatg cccaccacca gactcctgg gacctccaca      420
gtggatgtgg gaacctcagg gactccatcc tcacgcccc gcccccacg      468

```

```
<210> 115
```

<211> 468

<212> DNA

<213> Homo sapiens

<400> 115

```
actgctggcc ctctcctgat accattcacc ctcaacttca ccatcaccaa cctgcagtat    60
ggggaggaca tgggtcacc ttgctccagg aagttcaaca ccacagagag ggtcctgcag    120
ggtctgcttg gtcccatatt caagaacacc agtgttgccc ctctgtactc tggctgcaga    180
ctgacctctc tcaggctcga gaaggatgga gcagccactg gagtggatgc catctgcata    240
catcatottg accccaaaag ccttggactc aacagagagc ggctgtactg ggagctgagc    300
caactgacca atggcatcaa agagctgggc ccctacaccc tggacaggaa cagtctctat    360
gtcaatggtt tcacccatcg gacctctgtg cccaccacca gcaactcctg gacctccaca    420
gtggaccttg gaacctcagg gactccattc tcctcccaa gccccgca    468
```

<210> 116

<211> 468

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)..(468)

<223> All N's = any nucleotide

<400> 116

```
actgctggcc ctctcctggt gctgttcacc ctcaacttca ccatcaccaa cctgaagtat    60
gaggaggaca tgcctgcgcc ttgctccagg aagttcaaca ccaactgagag ggtcctgcag    120
actctgcttg gtctatggt caagaacacc agtgttgccc ttctgtactc tggctgcaga    180
ctgaccttgc tcaggctcga gaaggatgga gcagccactg gagtggatgc catctgcacc    240
caccgtcttg accccaaaag ccttggactg nacagngagc ngctntactg ggagctnagc    300
canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat    360
```

```
gtcaatggtt tcaccatcn ganctctgng cccaccacca gactcctgg gacctccaca 420
gtgnaentng gnaactcngg gactccatcc tcntccccc gccncaca 468
```

```
<210> 117
```

```
<211> 468
```

```
<212> DNA
```

```
<213> Homo sapiens
```

```
<220>
```

```
<221> misc_feature
```

```
<222> (1)..(468)
```

```
<223> All N's = any nucleotide
```

```
<400> 117
nennctgncc ctctcctgnt ncnnttcacc ntcaacttna ccatcaccaa cctgcantan 60
gnggannaca tgcnnncccc nggntccagg aagttcaaca ccacngagag agtccttcag 120
ggtctgctca ggccctgtgt caagaacacc agtggtggcc ctctgtactc tggctgcaga 180
ctgaccttgc tcaggcccaa gaaggatggg gcagccacca aagtggatgc catctgcacc 240
taccgccctg atcccaaaag cctgggactg gacagagagc agctatactg ggagctgagc 300
cagctaacc cagacatcac tgagctgggc ccctacacc aggacaggga cagtctctat 360
gtcaatggct tcaccatcg gagctctgtg ccaaccacca gtattcctgg gacctctgca 420
gtgcacctgg aaaccactgg gactccatcc tccttcccc gccacaca 468
```

```
<210> 118
```

```
<211> 468
```

```
<212> DNA
```

```
<213> Homo sapiens
```

```
<400> 118
gagcctggcc ctctcctgat accattcact ttcaacttta ccatcaccaa cctgcgttat 60
gaggaaaaca tgcaacacc cgggtccagg aagttcaaca ccacggagag ggttctgcag 120
ggtctgctca cgccctgtgt caagaacacc agtggtggcc ctctgtactc tggctgcaga 180
```

```

ctgaccttgc tcagacctga gaagcaggag gcagccactg gagtggacac catctgtacc 240
caccgcgttg atcccatcgg acctggactg gacagagagc ggctatactg ggagctgagc 300
cagctgacca acagcatcac agagctggga ccctacaccc tggataggga cagtctctat 360
gtgatggct tcaacccttg gagctctgtg ccaaccacca gcactcctgg gacctccaca 420
gtgcacctgg caacctctgg gactccatcc cccctgcctg gccacaca 468

```

<210> 119

<211> 468

<212> DNA

<213> Homo sapiens

```

<400> 119
gccccgtgcc ctctcttgat accattcacc ctcaacttta ccacaccga cctgcattat 60
gaagaaaaca tgcaacaccc tggttccagg aagtccaaca ccacggagag ggtctgcag 120
ggtctgctca agcccttggt caagagcacc agcgttggcc ctctgtactc tggctgcaga 180
ctgaccttgc tcagacctga gaaacatggg gcagccactg gagtggacgc catctgcacc 240
ctccgccttg atcccaactg tccctggactg gacagagagc ggctatactg ggagctgagc 300
cagctgacca acagcatcac agagctggga ccctacaccc tggataggga cagtctctat 360
gtcaatggct tcaacccttg gagctctgtg ccaaccacca gcactcctgg gacctccaca 420
gtgcacctgg caacctctgg gactccatcc tccctgcctg gccacaca 468

```

<210> 120

<211> 468

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)..(468)

<223> All N's = any nucleotide


```

<400> 120
actgctggcc ctctcctggt gccgttcacc ctcaacttca ccacaccaa cctgaagtac      60
gaggaggaca tgcattgccc tggctccagg aagtccaaca ccacagagag agtctcgacg      120
agtctgcatg gtcccatggt caagaacacc agtggtggcc ctctgtactc tggctgcaga      180
ctgaccttgc tcaggtcoga gaaggatgga gcagccactg gagtggatgc catctgcacc      240
caccgtcttg accccaaaag cctgggactg nacagnagac ngetntactg ggagctnagc      300
canctgacca annncatcnn ngagctgggn cctacacccc tggacaggna cagtctctat      360
gtcaatggtt tcacccatcn ganctctgng ccacacacca gactcctggt gacctccaca      420
gtgnaentng gnacctcngg gactccatcc tccntccccc gccncaca      468

```

```
<210> 121
```

```
<211> 468
```

```
<212> DNA
```

```
<213> Homo sapiens
```

```
<220>
```

```
<221> misc_feature
```

```
<222> (1)..(468)
```

```
<223> All N's = any nucleotide
```

```

<400> 121
nccnctgncc ctctcctgnt nccnttcacc ntcaacttna ccacaccaa cctgcantan      60
gnggannaca tgcnnncccc nggntccagg aagtccaaca ccacngagng ngtnctgcag      120
ggtctgtcnn ncccntntt caagaacccc agtgtnggcc ntctgtactc tggctgcaga      180
ctgacctnnc tcaggncnga gaagnatggn gcagccactg gantggatgc catctgcanc      240
caccnncntn ancccaaaag nccctggactg nacagnagac ngetntactg ggagctnagc      300
canctgacca acagcatcac agagctggga cctacacccc tggataggga cagtctctat      360
gtcaatggtt tcacccatcg aagctctatg ccacacacca gtattcctgg gacctctgca      420
gtgcacctgg aaacctctgg gactccagcc tccctccctg gccacaca      468

```

<210> 122
 <211> 468
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)..(468)
 <223> All N's = any nucleotide

<400> 122
 gccccctggcc ctctcctggt gccattcacc ctcaacttca ctatcaccaa cctgcagtat 60
 gaggaggaca tgcgtcacc tggttccagg aagttcaaca ccacggagag agtcctgcag 120
 ggtctgctca agcccttggt caagagcacc agtggtggcc ctctgtactc tggctgcaga 180
 ctgacccctgc tcaggcctga aaaacgtggg gcagccaccg gcgtggacac catctgcact 240
 caccgccttg accctctaaa ccctggactg nacagngagc ngctntactg ggagctnagc 300
 canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat 360
 gtcaatgggt tcacccatcn ganctctgng ccacaccaca gcactcctgg gacctccaca 420
 gtgnacntng gnacctcng gactccatcc tcntcccn gccncaca 468

<210> 123
 <211> 468
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)..(468)
 <223> All N's = any nucleotide

<400> 123

```

nennctgncc ctctcctgnt nocnttcacc ntcaacttna ccatcaccaa cctgcantn      60
gnggannaca tgcnnncccc nggntccagg aagttcaaca ccacngagng ngtnctgcag      120
ggtctgctnn nccccntntt caagaacncc agtgtnggcc ntctgtactc tggctgcaga      180
ctgacctnnc tcaggncnga gaagnatggn gcagccactg gantggatgc catctgcanc      240
caccnntnnt ancccaaaag nccctggactg nacagngagc ngctntactg ggagctnagc      300
canctgacca annncatenn ngagctgggn ccctacaccc tggacaggna cagtctctat      360
gtcaatgggt ttcacctctg gagctctgtg ccaaccacca gcaactcctgg gacctccaca      420
gtgcacctgg caacctctgg gactccatcc tcctgcctgg gccacaca      468

```

<210> 124

<211> 468

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)..(468)

<223> All N's = any nucleotide

```

<400> 124
gccctgtctc ctctcttgat accattcacc ctcaacttta ccatcaccaa cctgcattat      60
gaagaaaaca tgcaacaccc tggttccagg aagttcaaca ccacggagcg ggtctgcag      120
ggtctgcttg gtcccatgtt caagaacaca agtgtcggcc ttctgtactc tggctgcaga      180
ctgaccttgc tcaggcctga gaagaatggg gcagccactg gaatggatgc catctgcagc      240
caccgtcttg accccaaaag cccctggactg nacagngagc ngctntactg ggagctnagc      300
canctgacca annncatenn ngagctgggn ccctacaccc tggacaggna cagtctctat      360
gtcaatgggt tcaccatctn ganctctgng cccaccacca gcaactcctgg gacctccaca      420
gtgnacntng gnacctcngg gactccatcc tcctccccc gccncaca      468

```

<210> 125

<211> 468

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)..(468)

<223> All N's = any nucleotide

```

<400> 125
nccnctgncc ctctcctgnt nccnttcacc ntcaacttna ccatcaccaa cctgcantan    60
gnggannaca tgcnnncccc nggntccagg aagttcaaca ccacngagng ngtnctgcag    120
ggtctgctnn nccccntntt caagaacncc agtgtnggcc ntctgtactc tggctgcaga    180
ctgacctnnc tcaggnonga gaagnatggn gcagccactg gantggatgc catctgcanc    240
caccnnctnn ancccaaaag nccctggactg nacagnagac ngctntactg ggagctnagc    300
canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat    360
gtcaatggtt tcacccatca gaactctgtg ccacaccaca gtactcctgg gacctccaca    420
gtgtactggg caacctactg gactccatcc tcttccccg gccacaca                    468

```

<210> 126

<211> 468

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)..(468)

<223> All N's = any nucleotide

```

<400> 126
gagcctggcc ctctcctgat accattcact ttcaacttta ccatcaccaa cctgcattat    60
gaggaataca tgcaacaccc tggttccagg aagttcaaca ccacggagag ggttctgcag    120

```

```

ggctctgctca cgccctgtgt caagaacacc agtgttggcc ctctgtactc tggctgcaga      180
ctgaccttgc tcagacctga gaagcaggag gcagccactg gaggggacac catctgtacc      240
caccgcgttg atcccatcgg acctggactg nacagngagc ngctntactg ggagctnagc      300
canctgacca annncatenn ngagctgggn cctacacccc tggacaggna cagtctctat      360
gtcaatgggt tcacctaten ganctctgng cccaccacca gcactctctg gacctocaca      420
gtgnacntng gnacctcngg gactccatcc tccntccccc gccncaca      468

```

<210> 127

<211> 468

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)..(468)

<223> All N's = any nucleotide

```

<400> 127
nennctgncc ctctcctgnt nccnttcacc ntcaacttna ccatcaccaa cctgcantan      60
gnggannaca tgcnnncccc nggntccagg aagtccaaca ccacngagng ngtnctgcag      120
ggctctgctnn nncccntntt caagaacncc agtgtnggcc ntctgtactc tggctgcaga      180
ctgaccttnc tcaggncnga gaagnatggn gcagccactg gantggatgc catctgcanc      240
caccnncntn ancccaaaag nccctggactg nacagngagc ngctntactg ggagctnagc      300
canctgacca annncatenn ngagctgggn cctacacccc tggacaggna cagtctctat      360
gtcaatgggt tcacctatcg gagctctgtg ccaaccacca gcagctcttg gacctocaca      420
gtgcacctgg caacctctgg gactccatcc tccctgcttg gccacaca      468

```

<210> 128

<211> 468

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)..(468)

<223> All N's = any nucleotide

```
<400> 128
gccccgtgcc ctctcttgat accattcacc ctcaacttta ccatcaccaa cctgcattat    60
gaagaaaaca tgcaacaccc tggttccagg aagtccaaca ccacggagag ggttctgcag    120
ggtctgctca agcccttggt caagagcacc agtggtggcc ctctgtactc tggtctgcaga    180
ctgaccttgc tcagacctga gaaacatggg gcagccactg gagtggacgc catctgcacc    240
ctccgcttgg atcccactgg tctctggactg nacagnagag ngtntactg ggagctnagc    300
canctgacca annncatcnn ngagctgggn cctacacccc tggacaggna cagtctctat    360
gtcaatggtt tcaccatcnn ganctctgng cccaccacca gactcctgag gacctccaca    420
gtgnaentng gnacctcngg gactccatcc tcntccccn gcncacaa    468
```

<210> 129

<211> 468

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)..(468)

<223> All N's = any nucleotide

```
<400> 129
nennctgncc ctctcctgnt ncnnttcacc ntcaacttna ccatcaccaa cctgcantan    60
gnggannaca tgcnncccc nggntccagg aagtccaaca ccacngagng ngtnctgcag    120
ggtctgctnn nncnctnttt caagaacncc agtgtnggcc ntctgtactc tggtctgcaga    180
ctgacctnnc tcaggncnga gaagnatggn gcagccactg gantggatgc catctgcanc    240
```

```

cacnnntn ancccaaaag nctggactg nacagngagc ngetntactg ggagctnagc 300
canctgacca annncatenn ngagctgggn ccctacaccc tggacagzna cagtctctat 360
gtcaatgggt tcacccatcg gacctctgtg cccaccacca gactcctgg gacctccaca 420
gtgcacctgg caacctctgg gactccatcc tccctgctg gccacaca 468

```

<210> 130

<211> 468

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)..(468)

<223> All N's = any nucleotide

```

<400> 130
gcccctgtcc ctctcttgat accattcacc ctcaacttta ccatcacaa cctgcagtat 60
gaggaggaca tgcctcgccc tggatctagg aagtccaaca ccacagagag ggtcctgcag 120
ggtctgctta gtccatttt caagaactcc agtggtggcc ctctgtactc tggctgcaga 180
ctgacctctc tcaggcccca gaaggatggg gcagcaactg gaatggatgc tgtctgcctc 240
taccacccta atcccaaaag acctggactg nacagngagc ngetntactg ggagctnagc 300
canctgacca annncatenn ngagctgggn ccctacaccc tggacagzna cagtctctat 360
gtcaatgggt tcacccatcn ganctctgng cccaccacca gactcctgg gacctccaca 420
gtgnaatngn gnacctngg gactccatcc tccntcccn gccacaca 468

```

<210> 131

<211> 468

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)..(468)

<223> All N's = any nucleotide

```
<400> 131
nennctgncc ctctcctgnt nocnttcacc ntcaacttna ccacaccaa cctgcantan    60
gnggannaca tgennncccc nggntccagg aagttcaaca ccacngagng ngtnctgcag    120
ggtctgctnn nccccntntt caagaacncc agtgtnggcc ntctgtactc tggctgcaga    180
ctgacctnnc tcaggncnga gaagnatggn gcagccactg gantggatgc catctgcanc    240
caccnncntn ancccaaaag nccctggactg nacagnagc ngctntactg ggagctnagc    300
canctgacca annncatenn ngagctgggn ccctacaccc tggacaggna cagtctctat    360
gtcaatggtt tcaccattg gagctctggg ctcaccacca gcactccttg gacttcaca    420
gttgaccttg gaacctcagg gactccatcc cccgtcccca gccccaca    468
```

<210> 132

<211> 468

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)..(468)

<223> All N's = any nucleotide

```
<400> 132
actgctggcc ctctcctggt gccattcacc etaaacttca ccacaccaa cctgcagtat    60
gaggaggaca tgcatcgccc tggatctagg aagttcaacg ccacagagag ggtcctgcag    120
ggtctgetta gtcccatatt caagaacacc agtggtggcc ctctgtactc tggctgcaga    180
ctgaccttgc tcagacctga gaagcaggag gcagccactg gagtggacac catctgtacc    240
caccgctgtg atcccatcgg acctggactg nacagnagc ngctntactg ggagctnagc    300
canctgacca annncatenn ngagctgggn ccctacaccc tggacaggna cagtctctat    360
```



```
gtcaatgggt tcaccatcn ganctctgng cccaccacca gactctctgg gacctccaca 420
gtgnaentng gnacctengg gactccatcc tcntccccc gccncaca 468
```

<210> 133

<211> 468

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)..(468)

<223> All N's = any nucleotide

```
<400> 133
nncnctgncc ctctcctgnt nccnttcacc ntcaacttna ccatcaccaa cctgcantan 60
gnggannaca tgcnnncccc nggntccagg aagttcaaca ccacngagng ngtnctgcag 120
ggtctgctnn nccccnttt caagaacncc agtgtnggcc ntctgtaetc tggctgcaga 180
ctgacctnnc tcaggncnga gaagnatggn gcagccactg gantggatgc catctgcanc 240
caccnncntn ancccaaaag ncctggactg nacagngagc ngctntactg ggagctnagc 300
canctgacca annncatcnn ngagctgggn ccctacacco tggacaggna cagtctctat 360
gtcaatgggt tcaccatcg gagctttggg ctcaccacca gactccttg gacttcoca 420
gttgaccttg gaacctcagg gactccatcc cccgtcccca gccccaca 468
```

<210> 134

<211> 468

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)..(468)

<223> All N's = any nucleotide

```
<400> 134
actgctggcc ctctcctggt gccattcacc ctaaacttca ccatacacc aa cctgcagtat      60
gaggaggaca tgcacgccc tggctccagg aagttcaaca ccacggagag ggtccttcag      120
ggtctgctta cgcccttgtt caggaacacc agtgtcagct ctctgtactc tggttgcaga      180
ctgaccttgc tcaggcctga gaaggatggg gcagccacca gagtggatgc tgtctgcacc      240
catcgtcctg accccaaaag ccctggactg nacagngagc ngctntactg ggagctnagc      300
canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat      360
gtcaatgggt tcacccatcn gantctgng cccaccacca gcactcctgg gacctccaca      420
gtgnacntng gnacctengg gactccatcc tccttccccn gccncaca      468
```

<210> 135

<211> 465

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)..(465)

<223> All N's = any nucleotide

```
<400> 135
nccnctgncc ctctcctgnt nccnttcacc ntcaacttna ccatacacc aa cctgcantan      60
gnngannaca tgcnncccc nggntccagg aagttcaaca ccacggagng ngtnctgcag      120
ggtctgctnn nccccntntt caagaacncc agtgtnggcc ntctgtactc tggctgcaga      180
ctgacctnnc tcaggncnga gaagnatggn gcagccactg gantggatgc catctgcanc      240
caccnncntn ancccaaaag nccctggactg nacagngagc ngctntactg ggagctnagc      300
canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat      360
gtcaatgggt tcacccattg gatccctgtg cccaccagca gcactcctgg gacctccaca      420
gtggaccttg ggtcagggac tccatccctc cccccagcc ccaca      465
```

<210> 136
 <211> 468
 <212> DNA
 <213> Homo sapiens

<220>

<221> misc_feature

<222> (1)..(468)

<223> All N's = any nucleotide

<400> 136
 actgctggcc ctctcctggg accattcacc ctcaacttca ccatacacc cctgcagtat 60
 ggggaggaca tgggtcacc tggctccagg aagtccaaca ccacagagag ggtcctgcag 120
 ggtctgcttg gtcccatatt caagaacacc agtggtggcc ctctgtactc tggctgcaga 180
 ctgacctctc tcaggtcoga gaaggatgga gcagccactg gagtggatgc catctgcatac 240
 catcatcttg accccaaaag ccctggactg nacagnagac ngctntactg ggagctnagc 300
 canctgacca annncatcnn ngagctgggn ccctacaccc tggacagagna cagtctctat 360
 gtcaatggtt tcacccatcn ganctctgng ccacaccaca gcactcctgg gacctccaca 420
 gtgnacntng gnacctcngg gactccatcc tcntcccn gccncaca 468

<210> 137
 <211> 468
 <212> DNA
 <213> Homo sapiens

<220>

<221> misc_feature

<222> (1)..(468)

<223> All N's = any nucleotide

```

<400> 137
nennctgncc ctctcctgnt ncnnttcacc ntcaacttna ccatcaccaa cctgcantan    60
gnggannaca tgcnnncccc nggntccagg aagttcaaca ccacngagng ngtnctgcag    120
ggtctgctnn nnnccntntt caagaacncc agtgtnggcc ntctgtactc tggctgcaga    180
ctgacctnnc tcaggncnga gaagnatggn gcagccactg gantggatgc catctgcanc    240
caccnnctnn ancccaaaag noctggactg nacagnagag ngctntactg ggagctnagc    300
canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat    360
gtcaatgggt tcacccatca gacctttgcg cccaacacca gcactcctgg gacctccaca    420
gtggaccttg ggacctcagg gactccatcc tccctcccca gccctaca                    468

```

```
<210> 138
```

```
<211> 468
```

```
<212> DNA
```

```
<213> Homo sapiens
```

```
<220>
```

```
<221> misc_feature
```

```
<222> (1)..(468)
```

```
<223> All N's = any nucleotide
```

```

<400> 138
tctgctggcc ctctcctggt gccattcacc ctcaacttca ccatcaccaa cctgcagtac    60
gaggaggaca tgcacacccc aggctccagg aagttcaaca ccacggagcg ggtcctgcag    120
ggtctgcttg gtccatggtt caagaacacc agtgctggcc ttctgtactc tggctgcaga    180
ctgaccttgc tcaggcctga gaagaatggg gcagccacca gagtggatgc tgtctgcacc    240
catcgtcctg accccaaaag ccttggaactg nacagnagag ngctntactg ggagctnagc    300
canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat    360
gtcaatgggt tcacccatcn ganctctgng ccacacacca gcactcctgg gacctccaca    420
gtgnaentng gnaectcngg gactccatcc tccntcccn gccncaca                    468

```

```
<210> 139
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<211> 468
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)..(468)
 <223> All N's = any nucleotide

<400> 139
 ncnnetgncc ctctcctgnt ncnnttcacc ntcaacttna ccatacacc aa cctgcantaa 60
 gnggannaca tgcnnncccc ngntccagg aagttcaaca ccacngagag ggtctgcag 120
 ggtctgctca agcccttggt caagagcacc agtggtggcc ctctgtattc tggctgcaga 180
 ctgaccttgc tcaggcctga gaaggacgga gtagccacca gagtggacgc catctgcacc 240
 caccgccctg accccaaaat ccttgggcta gacagacagc agctatactg ggagctgagc 300
 cagctgaccc acagcatcac tgagctggga ccctacacc tggatagggg cagtctctat 360
 gtcaatgggt tcaccagcg gagctctgtg ccaccacca gcactcctgg gactttcaca 420
 gtacagccgg aaacctctga gactccatca tccctccctg gcccacaa 468

<210> 140
 <211> 468
 <212> DNA
 <213> Homo sapiens

<400> 140
 gccactggcc ctgtcctgct gccattcacc ctcaatttta ccactactaa cctgcagtat 60
 gaggaggaca tgcctgcgcc tggctccagg aagttcaaca ccacggagag ggtccttcag 120
 ggtctgctta tgcccttggt caagaacacc agtgctcagct ctctgtactc tggctgcaga 180
 ctgaccttgc tcaggcctga gaaggatggg gcagccacca gagtggatgc tgtctgcacc 240
 catcgtcctg accccaaaag ccttggactg gacagagagc ggctgtactg gaagctgagc 300
 cagctgaccc acggcatcac tgagctgggc ccctacacc tggacaggca cagtctctat 360

```
gtcaatggtt tcacccatca gagctctatg acgaccacca gaactcctga tacctccaca 420
atgcacctgg caacctcgag aactccagcc tccctgtctg gacctacg 468
```

<210> 141

<211> 468

<212> DNA

<213> Homo sapiens

```
<400> 141
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gaggagaaca tgcattaccc tggtctctaga aagtttaaca ccacggagag agtccttcag 120
ggtctgctca ggctgtggtt caagaacacc agtggtggcc ctctgtactc tggctgcaga 180
ctgaccttgc tcaggcccaa gaaggatggg gcagccacca aagtggatgc catctgcacc 240
taccgccctg atcccaaaaag cctgggactg gacagagagc agctatactg ggagctgagc 300
cagctaacc cagacatcac tgagctgggc cctacacccc tggacaggga cagctcttat 360
gtcaatggtt tcacacagcg gagctctgtg cccaccacta gcattcctgg gacccccaca 420
gtggacctgg gaacatctgg gactccagtt tctaaacctg gtccctcg 468
```

<210> 142

<211> 468

<212> DNA

<213> Homo sapiens

```
<400> 142
gctgccagcc ctctcctggt gctattcact ctcaacttca ccaccaccaa cctgcgggtat 60
gaggagaaca tgcagcacc tggtccaggg aagttcaaca ccacggagag ggtccttcag 120
ggcctgctca ggtcctggtt caagagcacc agtggtggcc ctctgtactc tggctgcaga 180
ctgactttgc tcaggcctga aaaggatggg acagccactg gagtggatgc catctgcacc 240
caccaccctg accccaaaag cctaggctg gacagagagc agctgtattg ggagctgagc 300
cagctgaccc acaatatcac tgagctgggc cactatgccc tggacaacga cagcctcttt 360
gtcaatggtt tcaactcatg gagctctgtg tccaccacca gactcctgg gacccccaca 420
gtgtatctgg gagcatctaa gactccagcc tcgatatttg gcccttca 468
```

<210> 143

<211> 399

<212> DNA

<213> Homo sapiens

```

<400> 143
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gaggagaaca tgtggcctgg ctccaggaag ttcaacacta cagagagggg ccttcagggg    120
ctgctaaggc ccttggtcaa gaacaccagt gttggccctc tgtactctgg ctccaggctg    180
accttgctca ggccagagaa agatggggaa gccaccggag tggatgccat ctgacccac    240
cgccctgacc ccacaggccc tgggctggac agagagcagc tgtatttga gctgagccag    300
ctgaccaca gcatcactga gctgggcccc tacacactgg acagggacag tctctatgtc    360
aatggtttca cccatcggag ctctgtacco accaccagc                                399

```

<210> 144

<211> 453

<212> DNA

<213> Homo sapiens

```

<400> 144
accggggtgg tcagcgagga gccattcaca ctgaacttca ccatcaacaa cctgcgctac    60
atggcgagaa tgggccaacc cggtccctc aagttcaaca tcacagacaa cgctatgaag    120
cacctgctca gtcctttgtt ccagaggagc agcctgggtg cacggtacac aggctgcagg    180
gtcatcgcac taaggtctgt gaagaacggt gctgagacac ggggtggacct cctctgcacc    240
tacctgcagc ccctcagcgg ccaggtctg cctatcaagc aggtgttcca tgagctgagc    300
cagcagaccc atggcatcac ccgctggggc cctactctc tggacaaaga cagcctctac    360
cttaacggtt acaatgaacc tggctatagat gagectcta caactcccaa gccagccacc    420
acattctcgc ctctctctgc agaagccaca aca                                453

```

<210> 145

<211> 465

<212> DNA

<213> Homo sapiens

```

<400> 145
gccatgggggt accacctgaa gaccctcaca ctcaacttca ccactctcaa tctccagtat    60
tcaccagata tgggcaaggg ctgagctaca ttcaactcca ccgaggggggt ccttcagcac    120
ctgctcagac ccttgttcca gaagagcagc atgggcccct tctacttggg ttgccaaactg    180
atctccctca ggctgagaa ggatggggca gccactggtg tggacaccac ctgcacctac    240
caccctgacc ctgtggggccc cgggctggac atacagcagc ttacttggga gctgagtcag    300
ctgacccatg gtgtcaccca actgggcttc tatgtcctgg acagggatag cctcttcac    360
aatggctatg caccocagaa tttatcaatc cggggcgcagt accagataaa ttccacatt    420
gtcaactgga acctcagtaa tcagacccc acatcctcag agtac                                465

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<210> 146

<211> 9799

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (1)..(9799)

<223> Any "X" = any amino acid

<400> 146

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Ala Thr Val Pro Phe Met Val Pro Phe Thr Leu Asn Phe Thr Ile Thr
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Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe
          20          25          30
Asn Ala Thr Glu Arg Glu Leu Gln Gly Leu Leu Lys Pro Leu Phe Arg
          35          40          45
Asn Ser Ser Leu Glu Tyr Leu Tyr Ser Gly Cys Arg Leu Ala Ser Leu
          50          55          60

```


Arg Pro Glu Lys Asp Ser Ser Ala Met Ala Val Asp Ala Ile Cys Thr
 65 70 75 80
 His Arg Pro Asp Pro Glu Asp Leu Gly Leu Asp Arg Glu Arg Leu Tyr
 85 90 95
 Trp Glu Leu Ser Asn Leu Thr Asn Gly Ile Gln Glu Leu Gly Pro Tyr
 100 105 110
 Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser
 115 120 125
 Ser Met Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Val Gly
 130 135 140
 Thr Ser Gly Thr Pro Ser Ser Ser Pro Ser Pro Thr Ala Ala Gly Pro
 145 150 155 160
 Leu Leu Met Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr
 165 170 175
 Glu Glu Asp Met Arg Arg Thr Gly Ser Arg Lys Phe Asn Thr Met Glu
 180 185 190
 Ser Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val
 195 200 205
 Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys
 210 215 220
 Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu Asp
 225 230 235 240
 Pro Lys Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp Glu Leu Ser
 245 250 255
 Lys Leu Thr Asn Asp Ile Glu Glu Leu Gly Pro Tyr Thr Leu Asp Arg
 260 265 270
 Asn Ser Leu Tyr Val Asn Gly Phe Thr His Gln Ser Ser Val Ser Thr
 275 280 285
 Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Arg Thr Ser Gly Thr
 290 295 300
 Pro Ser Ser Leu Ser Ser Pro Thr Ile Met Ala Ala Gly Pro Leu Leu
 305 310 315 320
 Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Gly Glu
 325 330 335
 Asp Met Gly His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val
 340 345 350
 Leu Gln Gly Leu Leu Gly Pro Ile Phe Lys Asn Thr Ser Val Gly Pro
 355 360 365
 Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Ser Glu Lys Asp Gly

370					375					380					
Ala	Ala	Thr	Gly	Val	Asp	Ala	Ile	Cys	Ile	His	His	Leu	Asp	Pro	Lys
385					390					395					400
Ser	Pro	Gly	Leu	Asn	Arg	Glu	Arg	Leu	Tyr	Trp	Glu	Leu	Ser	Gln	Leu
				405					410					415	
Thr	Asn	Gly	Ile	Lys	Glu	Leu	Gly	Pro	Tyr	Thr	Leu	Asp	Arg	Asn	Ser
			420					425					430		
Leu	Tyr	Val	Asn	Gly	Phe	Thr	His	Arg	Thr	Ser	Val	Pro	Thr	Ser	Ser
			435				440					445			
Thr	Pro	Gly	Thr	Ser	Thr	Val	Asp	Leu	Gly	Thr	Ser	Gly	Thr	Pro	Phe
			450				455					460			
Ser	Leu	Pro	Ser	Pro	Ala	Thr	Ala	Gly	Pro	Leu	Leu	Val	Leu	Phe	Thr
					470					475					480
Leu	Asn	Phe	Thr	Ile	Thr	Asn	Leu	Lys	Tyr	Glu	Glu	Asp	Met	His	Arg
				485					490					495	
Pro	Gly	Ser	Arg	Lys	Phe	Asn	Thr	Thr	Glu	Arg	Val	Leu	Gln	Thr	Leu
			500					505					510		
Leu	Gly	Pro	Met	Phe	Lys	Asn	Thr	Ser	Val	Gly	Leu	Leu	Tyr	Ser	Gly
			515					520					525		
Cys	Arg	Leu	Thr	Leu	Leu	Arg	Ser	Glu	Lys	Asp	Gly	Ala	Ala	Thr	Gly
			530				535				540				
Val	Asp	Ala	Ile	Cys	Thr	His	Arg	Leu	Asp	Pro	Lys	Ser	Pro	Gly	Leu
			545			550				555					560
Asp	Arg	Glu	Gln	Leu	Tyr	Trp	Glu	Leu	Ser	Gln	Leu	Thr	Asn	Gly	Ile
			565						570					575	
Lys	Glu	Leu	Gly	Pro	Tyr	Thr	Leu	Asp	Arg	Asn	Ser	Leu	Tyr	Val	Asn
			580					585					590		
Gly	Phe	Thr	His	Trp	Ile	Pro	Val	Pro	Thr	Ser	Ser	Thr	Pro	Gly	Thr
			595				600						605		
Ser	Thr	Val	Asp	Leu	Gly	Ser	Gly	Thr	Pro	Ser	Ser	Leu	Pro	Ser	Pro
			610				615					620			
Thr	Ala	Ala	Gly	Pro	Leu	Leu	Val	Pro	Phe	Thr	Leu	Asn	Phe	Thr	Ile
			625			630					635				640
Thr	Asn	Leu	Gln	Tyr	Glu	Glu	Asp	Met	His	His	Pro	Gly	Ser	Arg	Lys
			645						650					655	
Phe	Asn	Thr	Thr	Glu	Arg	Val	Leu	Gln	Gly	Leu	Leu	Gly	Pro	Met	Phe
			660						665				670		
Lys	Asn	Thr	Ser	Val	Gly	Leu	Leu	Tyr	Ser	Gly	Cys	Arg	Leu	Thr	Leu
			675				680								

Leu Arg Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys
 690 695
 Thr His Arg Leu Asp Pro Lys Ser Pro Gly Val Asp Arg Glu Gln Leu
 705 710 715 720
 Tyr Trp Glu Leu Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly Pro
 725 730 735
 Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Gln
 740 745 750
 Thr Ser Ala Pro Asn Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu
 755 760 765
 Gly Thr Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr Ser Ala Gly
 770 775 780
 Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln
 785 790 795 800
 Tyr Glu Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe Asn Thr Thr
 805 810 815
 Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser
 820 825 830
 Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Ser Glu
 835 840 845
 Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu
 850 855 860
 Asp Pro Lys Ser Pro Gly Val Asp Arg Glu Gln Leu Tyr Trp Glu Leu
 865 870 875 880
 Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp
 885 890 895
 Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Gln Thr Ser Ala Pro
 900 905 910
 Asn Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly
 915 920 925
 Thr Pro Ser Ser Leu Pro Ser Pro Thr Ser Ala Gly Pro Leu Leu Val
 930 935 940
 Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp
 945 950 955 960
 Met His His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu
 965 970 975
 Gln Gly Leu Leu Gly Pro Met Phe Lys Asn Thr Ser Val Gly Leu Leu
 980 985 990
 Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asn Gly Ala
 995 1000 1005

Ala Thr	Gly Met Asp Ala	Ile	Cys Ser His Arg	Leu	Asp Pro Lys
1010		1015		1020	
Ser Pro	Gly Leu Asn Arg	Glu	Gln Leu Tyr Trp	Glu	Leu Ser Gln
1025		1030		1035	
Leu Thr	His Gly Ile Lys	Glu	Leu Gly Pro Tyr	Thr	Leu Asp Arg
1040		1045		1050	
Asn Ser	Leu Tyr Val Asn	Gly	Phe Thr His Arg	Ser	Ser Val Ala
1055		1060		1065	
Pro Thr	Ser Thr Pro Gly	Thr	Ser Thr Val Asp	Leu	Gly Thr Ser
1070		1075		1080	
Gly Thr	Pro Ser Ser Leu	Pro	Ser Pro Thr Thr	Ala	Val Pro Leu
1085		1090		1095	
Leu Val	Pro Phe Thr Leu	Asn	Phe Thr Ile Thr	Asn	Leu Gln Tyr
1100		1105		1110	
Gly Glu	Asp Met Arg His	Pro	Gly Ser Arg Lys	Phe	Asn Thr Thr
1115		1120		1125	
Glu Arg	Val Leu Gln Gly	Leu	Leu Gly Pro Leu	Phe	Lys Asn Ser
1130		1135		1140	
Ser Val	Gly Pro Leu Tyr	Ser	Gly Cys Arg Leu	Ile	Ser Leu Arg
1145		1150		1155	
Ser Glu	Lys Asp Gly Ala	Ala	Thr Gly Val Asp	Ala	Ile Cys Thr
1160		1165		1170	
His His	Leu Asn Pro Gln	Ser	Pro Gly Leu Asp	Arg	Glu Gln Leu
1175		1180		1185	
Tyr Trp	Gln Leu Ser Gln	Met	Thr Asn Gly Ile	Lys	Glu Leu Gly
1190		1195		1200	
Pro Tyr	Thr Leu Asp Arg	Asn	Ser Leu Tyr Val	Asn	Gly Phe Thr
1205		1210		1215	
His Arg	Ser Ser Gly Leu	Thr	Thr Ser Thr Pro	Trp	Thr Ser Thr
1220		1225		1230	
Val Asp	Leu Gly Thr Ser	Gly	Thr Pro Ser Pro	Val	Pro Ser Pro
1235		1240		1245	
Thr Thr	Ala Gly Pro Leu	Leu	Val Pro Phe Thr	Leu	Asn Phe Thr
1250		1255		1260	
Ile Thr	Asn Leu Gln Tyr	Glu	Glu Asp Met His	Arg	Pro Gly Ser
1265		1270		1275	
Arg Lys	Phe Asn Ala Thr	Glu	Arg Val Leu Gln	Gly	Leu Leu Ser
1280		1285		1290	
Pro Ile	Phe Lys Asn Ser	Ser	Val Gly Pro Leu	Tyr	Ser Gly Cys

1295	1300	1305
Arg Leu Thr Ser Leu Arg	Pro Glu Lys Asp Gly Ala	Ala Thr Gly
1310	1315	1320
Met Asp Ala Val Cys Leu	Tyr His Pro Asn Pro Lys	Arg Pro Gly
1325	1330	1335
Leu Asp Arg Glu Gln Leu	Tyr Trp Glu Leu Ser Gln	Leu Thr His
1340	1345	1350
Asn Ile Thr Glu Leu Gly	Pro Tyr Ser Leu Asp Arg	Asp Ser Leu
1355	1360	1365
Tyr Val Asn Gly Phe Thr	His Gln Asn Ser Val Pro	Thr Thr Ser
1370	1375	1380
Thr Pro Gly Thr Ser Thr	Val Tyr Trp Ala Thr Thr	Gly Thr Pro
1385	1390	1395
Ser Ser Phe Pro Gly His	Thr Glu Pro Gly Pro Leu	Leu Ile Pro
1400	1405	1410
Phe Thr Phe Asn Phe Thr	Ile Thr Asn Leu His Tyr	Glu Glu Asn
1415	1420	1425
Met Gln His Pro Gly Ser	Arg Lys Phe Asn Thr Thr	Glu Arg Val
1430	1435	1440
Leu Gln Gly Leu Leu Lys	Pro Leu Phe Lys Asn Thr	Ser Val Gly
1445	1450	1455
Pro Leu Tyr Ser Gly Cys	Arg Leu Thr Ser Leu Arg	Pro Glu Lys
1460	1465	1470
Asp Gly Ala Ala Thr Gly	Met Asp Ala Val Cys Leu	Tyr His Pro
1475	1480	1485
Asn Pro Lys Arg Pro Gly	Leu Asp Arg Glu Gln Leu	Tyr Cys Glu
1490	1495	1500
Leu Ser Gln Leu Thr His	Asn Ile Thr Glu Leu Gly	Pro Tyr Ser
1505	1510	1515
Leu Asp Arg Asp Ser Leu	Tyr Val Asn Gly Phe Thr	His Gln Asn
1520	1525	1530
Ser Val Pro Thr Thr Ser	Thr Pro Gly Thr Ser Thr	Val Tyr Trp
1535	1540	1545
Ala Thr Thr Gly Thr Pro	Ser Ser Phe Pro Gly His	Thr Glu Pro
1550	1555	1560
Gly Pro Leu Leu Ile Pro	Phe Thr Phe Asn Phe Thr	Ile Thr Asn
1565	1570	1575
Leu His Tyr Glu Glu Asn	Met Gln His Pro Gly Ser	Arg Lys Phe
1580	1585	1590

Asn Thr	Thr Glu Arg Val	Leu	Gln Gly Leu Leu	Lys	Pro Leu Phe
1595		1600		1605	
Lys Asn	Thr Ser Val Gly	Pro	Leu Tyr Ser Gly	Cys	Arg Leu Thr
1610		1615		1620	
Leu Leu	Arg Pro Glu Lys	His	Glu Ala Ala Thr	Gly	Val Asp Thr
1625		1630		1635	
Ile Cys	Thr His Arg Val	Asp	Pro Ile Gly Pro	Gly	Leu Asp Arg
1640		1645		1650	
Glu Arg	Leu Tyr Trp Glu	Leu	Ser Gln Leu Thr	Asn	Ser Ile Thr
1655		1660		1665	
Glu Leu	Gly Pro Tyr Thr	Leu	Asp Arg Asp Ser	Leu	Tyr Val Asn
1670		1675		1680	
Gly Phe	Asn Pro Arg Ser	Ser	Val Pro Thr Thr	Ser	Thr Pro Gly
1685		1690		1695	
Thr Ser	Thr Val His Leu Ala	Thr	Ser Gly Thr Pro	Ser	Ser Leu
1700		1705		1710	
Pro Gly	His Thr Ala Pro	Val	Pro Leu Leu Ile	Pro	Phe Thr Leu
1715		1720		1725	
Asn Phe	Thr Ile Thr Asn	Leu	His Tyr Glu Glu	Asn	Met Gln His
1730		1735		1740	
Pro Gly	Ser Arg Lys Phe	Asn	Thr Thr Glu Arg	Val	Leu Gln Gly
1745		1750		1755	
Leu Leu	Lys Pro Leu Phe	Lys	Asn Thr Ser Val	Gly	Pro Leu Tyr
1760		1765		1770	
Ser Gly	Cys Arg Leu Thr	Leu	Leu Arg Pro Glu	Lys	His Glu Ala
1775		1780		1785	
Ala Thr	Gly Val Asp Thr	Ile	Cys Thr His Arg	Val	Asp Pro Ile
1790		1795		1800	
Gly Pro	Gly Leu Asp Arg	Glu	Xaa Leu Tyr Trp	Glu	Leu Ser Xaa
1805		1810		1815	
Leu Thr	Xaa Xaa Ile Xaa	Glu	Leu Gly Pro Tyr	Xaa	Leu Asp Arg
1820		1825		1830	
Xaa Ser	Leu Tyr Val Asn	Gly	Phe Xaa Xaa Xaa	Xaa	Xaa Xaa Xaa
1835		1840		1845	
Xaa Thr	Ser Thr Pro Gly	Thr	Ser Xaa Val Xaa	Leu	Xaa Thr Ser
1850		1855		1860	
Gly Thr	Pro Xaa Xaa Xaa	Pro	Xaa Xaa Thr Ser	Ala	Gly Pro Leu
1865		1870		1875	
Leu Val	Pro Phe Thr Leu	Asn	Phe Thr Ile Thr	Asn	Leu Gln Tyr
1880		1885		1890	

Glu	Glu	Asp	Met	His	His	Pro	Gly	Ser	Arg	Lys	Phe	Asn	Thr	Thr
1895						1900					1905			
Glu	Arg	Val	Leu	Gln	Gly	Leu	Leu	Gly	Pro	Met	Phe	Lys	Asn	Thr
1910						1915					1920			
Ser	Val	Gly	Leu	Leu	Tyr	Ser	Gly	Cys	Arg	Leu	Thr	Leu	Leu	Arg
1925						1930					1935			
Pro	Glu	Lys	Asn	Gly	Ala	Ala	Thr	Gly	Met	Asp	Ala	Ile	Cys	Ser
1940						1945					1950			
His	Arg	Leu	Asp	Pro	Lys	Ser	Pro	Gly	Leu	Asp	Arg	Glu	Gln	Leu
1955						1960					1965			
Tyr	Trp	Glu	Leu	Ser	Gln	Leu	Thr	His	Gly	Ile	Lys	Glu	Leu	Gly
1970						1975					1980			
Pro	Tyr	Thr	Leu	Asp	Arg	Asn	Ser	Leu	Tyr	Val	Asn	Gly	Phe	Thr
1985						1990					1995			
His	Arg	Ser	Ser	Val	Ala	Pro	Thr	Ser	Thr	Pro	Gly	Thr	Ser	Thr
2000						2005					2010			
Val	Asp	Leu	Gly	Thr	Ser	Gly	Thr	Pro	Ser	Ser	Leu	Pro	Ser	Pro
2015						2020					2025			
Thr	Thr	Ala	Val	Pro	Leu	Leu	Val	Pro	Phe	Thr	Leu	Asn	Phe	Thr
2030						2035					2040			
Ile	Thr	Asn	Leu	Gln	Tyr	Gly	Glu	Asp	Met	Arg	His	Pro	Gly	Ser
2045						2050					2055			
Arg	Lys	Phe	Asn	Thr	Thr	Glu	Arg	Val	Leu	Gln	Gly	Leu	Leu	Gly
2060						2065					2070			
Pro	Leu	Phe	Lys	Asn	Ser	Ser	Val	Gly	Pro	Leu	Tyr	Ser	Gly	Cys
2075						2080					2085			
Arg	Leu	Ile	Ser	Leu	Arg	Ser	Glu	Lys	Asp	Gly	Ala	Ala	Thr	Gly
2090						2095					2100			
Val	Asp	Ala	Ile	Cys	Thr	His	His	Leu	Asn	Pro	Gln	Ser	Pro	Gly
2105						2110					2115			
Leu	Asp	Arg	Glu	Gln	Leu	Tyr	Trp	Gln	Leu	Ser	Gln	Met	Thr	Asn
2120						2125					2130			
Gly	Ile	Lys	Glu	Leu	Gly	Pro	Tyr	Thr	Leu	Asp	Arg	Asn	Ser	Leu
2135						2140					2145			
Tyr	Val	Asn	Gly	Phe	Thr	His	Arg	Ser	Ser	Gly	Leu	Thr	Thr	Ser
2150						2155					2160			
Thr	Pro	Trp	Thr	Ser	Thr	Val	Asp	Leu	Gly	Thr	Ser	Gly	Thr	Pro
2165						2170					2175			
Ser	Pro	Val	Pro	Ser	Pro	Thr	Thr	Ala	Gly	Pro	Leu	Leu	Val	Pro

2180	2185	2190
Phe Thr Leu Asn Phe Thr	Ile Thr Asn Leu Gln Tyr	Glu Glu Asp
2195	2200	2205
Met His Arg Pro Gly Ser	Arg Lys Phe Asn Ala Thr	Glu Arg Val
2210	2215	2220
Leu Gln Gly Leu Leu Ser	Pro Ile Phe Lys Asn Ser	Ser Val Gly
2225	2230	2235
Pro Leu Tyr Ser Gly Cys	Arg Leu Thr Ser Leu Arg	Pro Glu Lys
2240	2245	2250
Asp Gly Ala Ala Thr Gly	Met Asp Ala Val Cys Leu	Tyr His Pro
2255	2260	2265
Asn Pro Lys Arg Pro Gly	Leu Asp Arg Glu Gln Leu	Tyr Trp Glu
2270	2275	2280
Leu Ser Gln Leu Thr His	Asn Ile Thr Glu Leu Gly	Pro Tyr Ser
2285	2290	2295
Leu Asp Arg Asp Ser Leu	Tyr Val Asn Gly Phe Thr	His Gln Ser
2300	2305	2310
Ser Met Thr Thr Thr Arg	Thr Pro Asp Thr Ser Thr	Met His Leu
2315	2320	2325
Ala Thr Ser Arg Thr Pro	Ala Ser Leu Ser Gly Pro	Thr Thr Ala
2330	2335	2340
Ser Pro Leu Leu Val Leu	Phe Thr Ile Asn Cys Thr	Ile Thr Asn
2345	2350	2355
Leu Gln Tyr Glu Glu Asp	Met Arg Arg Thr Gly Ser	Arg Lys Phe
2360	2365	2370
Asn Thr Met Glu Ser Val	Leu Gln Gly Leu Leu Lys	Pro Leu Phe
2375	2380	2385
Lys Asn Thr Ser Val Gly	Pro Leu Tyr Ser Gly Cys	Arg Leu Thr
2390	2395	2400
Leu Leu Arg Pro Lys Lys	Asp Gly Ala Ala Thr Gly	Val Asp Ala
2405	2410	2415
Ile Cys Thr His Arg Leu	Asp Pro Lys Ser Pro Gly	Leu Asn Arg
2420	2425	2430
Glu Gln Leu Tyr Trp Glu	Leu Ser Lys Leu Thr Asn	Asp Ile Glu
2435	2440	2445
Glu Leu Gly Pro Tyr Thr	Leu Asp Arg Asn Ser Leu	Tyr Val Asn
2450	2455	2460
Gly Phe Thr His Gln Ser	Ser Val Ser Thr Thr Ser	Thr Pro Gly
2465	2470	2475

Thr Ser	Thr Val Asp Leu Arg	Thr Ser Gly Thr Pro	Ser Ser Leu
2480	2485	2490	
Ser Ser	Pro Thr Ile Met Xaa	Xaa Xaa Pro Leu Leu	Xaa Pro Phe
2495	2500	2505	
Thr Leu	Asn Phe Thr Ile Thr	Asn Leu Xaa Tyr Glu	Glu Xaa Met
2510	2515	2520	
Xaa Xaa	Pro Gly Ser Arg Lys	Phe Asn Thr Thr Glu	Arg Val Leu
2525	2530	2535	
Gln Gly	Leu Leu Arg Pro Leu	Phe Lys Asn Thr Ser	Val Ser Ser
2540	2545	2550	
Leu Tyr	Ser Gly Cys Arg Leu	Thr Leu Leu Arg Pro	Glu Lys Asp
2555	2560	2565	
Gly Ala	Ala Thr Arg Val Asp	Ala Ala Cys Thr Tyr	Arg Pro Asp
2570	2575	2580	
Pro Lys	Ser Pro Gly Leu Asp	Arg Glu Gln Leu Tyr	Trp Glu Leu
2585	2590	2595	
Ser Gln	Leu Thr His Ser Ile	Thr Glu Leu Gly Pro	Tyr Thr Leu
2600	2605	2610	
Asp Arg	Val Ser Leu Tyr Val	Asn Gly Phe Asn Pro	Arg Ser Ser
2615	2620	2625	
Val Pro	Thr Thr Ser Thr Pro	Gly Thr Ser Thr Val	His Leu Ala
2630	2635	2640	
Thr Ser	Gly Thr Pro Ser Ser	Leu Pro Gly His Thr	Ala Pro Val
2645	2650	2655	
Pro Leu	Leu Ile Pro Phe Thr	Leu Asn Phe Thr Ile	Thr Asn Leu
2660	2665	2670	
His Tyr	Glu Glu Asn Met Gln	His Pro Gly Ser Arg	Lys Phe Asn
2675	2680	2685	
Thr Thr	Glu Arg Val Leu Gln	Gly Leu Leu Arg Pro	Leu Phe Lys
2690	2695	2700	
Ser Thr	Ser Val Gly Pro Leu	Tyr Ser Gly Cys Arg	Leu Thr Leu
2705	2710	2715	
Leu Arg	Pro Glu Lys His Gly	Ala Ala Thr Gly Val	Asp Ala Ile
2720	2725	2730	
Cys Thr	Leu Arg Leu Asp Pro	Thr Gly Pro Gly Leu	Asp Arg Glu
2735	2740	2745	
Arg Leu	Tyr Trp Glu Leu Ser	Gln Leu Thr Asn Ser	Val Thr Glu
2750	2755	2760	
Leu Gly	Pro Tyr Thr Leu Asp	Arg Asp Ser Leu Tyr	Val Asn Gly
2765	2770	2775	

Phe Thr Gln Arg Ser Ser Val Pro Thr Thr Ser Ile Pro Gly Thr
 2780 2785 2790
 Ser Ala Val His Leu Glu Thr Ser Gly Thr Pro Ala Ser Leu Pro
 2795 2800 2805
 Gly His Thr Ala Pro Gly Pro Leu Leu Val Pro Phe Thr Leu Asn
 2810 2815 2820
 Phe Thr Ile Thr Asn Leu Gln Tyr Glu Val Asp Met Arg His Pro
 2825 2830 2835
 Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu
 2840 2845 2850
 Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser
 2855 2860 2865
 Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Arg Gly Ala Ala
 2870 2875 2880
 Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu Asp Pro Leu Asn
 2885 2890 2895
 Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu
 2900 2905 2910
 Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp Arg Gly
 2915 2920 2925
 Ser Leu Tyr Val Asn Gly Phe Thr His Arg Asn Phe Val Pro Ile
 2930 2935 2940
 Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Gly Thr Ser Glu
 2945 2950 2955
 Thr Pro Ser Ser Leu Pro Arg Pro Ile Val Pro Gly Pro Leu Leu
 2960 2965 2970
 Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu
 2975 2980 2985
 Glu Ala Met Arg His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu
 2990 2995 3000
 Arg Val Leu Gln Gly Leu Leu Arg Pro Leu Phe Lys Asn Thr Ser
 3005 3010 3015
 Ile Gly Pro Leu Tyr Ser Ser Cys Arg Leu Thr Leu Leu Arg Pro
 3020 3025 3030
 Glu Lys Asp Lys Ala Ala Thr Arg Val Asp Ala Ile Cys Thr His
 3035 3040 3045
 His Pro Asp Pro Gln Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr
 3050 3055 3060
 Trp Glu Leu Ser Gln Leu Thr His Gly Ile Thr Glu Leu Gly Pro

3065	3070	3075
Tyr Thr Leu Asp Arg Asp Ser 3080	Leu Tyr Val Asp Gly 3085	Phe Thr His 3090
Trp Ser Pro Ile Pro Thr 3095	Thr Ser Thr Pro Gly 3100	Thr Ser Ile Val 3105
Asn Leu Gly Thr Ser Gly 3110	Ile Pro Pro Ser Leu 3115	Pro Glu Thr Thr 3120
Xaa Xaa Xaa Pro Leu Leu 3125	Xaa Pro Phe Thr Leu 3130	Asn Phe Thr Ile 3135
Thr Asn Leu Xaa Tyr Glu 3140	Glu Xaa Met Xaa Xaa 3145	Pro Gly Ser Arg 3150
Lys Phe Asn Thr Thr Glu 3155	Arg Val Leu Gln Gly 3160	Leu Leu Lys Pro 3165
Leu Phe Arg Asn Ser Ser 3170	Leu Glu Tyr Leu Tyr 3175	Ser Gly Cys Arg 3180
Leu Ala Ser Leu Arg Pro 3185	Glu Lys Asp Ser Ser 3190	Ala Met Ala Val 3195
Asp Ala Ile Cys Thr His 3200	Arg Pro Asp Pro Glu 3205	Asp Leu Gly Leu 3210
Asp Arg Glu Arg Leu Tyr 3215	Trp Glu Leu Ser Asn 3220	Leu Thr Asn Gly 3225
Ile Gln Glu Leu Gly Pro 3230	Tyr Thr Leu Asp Arg 3235	Asn Ser Leu Tyr 3240
Val Asn Gly Phe Thr His 3245	Arg Ser Ser Phe Leu 3250	Thr Thr Ser Thr 3255
Pro Trp Thr Ser Thr Val 3260	Asp Leu Gly Thr Ser 3265	Gly Thr Pro Ser 3270
Pro Val Pro Ser Pro Thr 3275	Thr Ala Gly Pro Leu 3280	Leu Val Pro Phe 3285
Thr Leu Asn Phe Thr Ile 3290	Thr Asn Leu Gln Tyr 3295	Glu Glu Asp Met 3300
His Arg Pro Gly Ser Arg 3305	Arg Phe Asn Thr Thr 3310	Glu Arg Val Leu 3315
Gln Gly Leu Leu Thr Pro 3320	Leu Phe Lys Asn Thr 3325	Ser Val Gly Pro 3330
Leu Tyr Ser Gly Cys Arg 3335	Leu Thr Leu Leu Arg 3340	Pro Glu Lys Gln 3345
Glu Ala Ala Thr Gly Val 3350	Asp Thr Ile Cys Thr 3355	His Arg Val Asp 3360

Pro Ile Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu
 3365 3370 3375
 Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu
 3380 3385 3390
 Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Asn Pro Trp Ser Ser
 3395 3400 3405
 Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala
 3410 3415 3420
 Thr Ser Gly Thr Pro Ser Ser Leu Pro Gly His Thr Ala Pro Val
 3425 3430 3435
 Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr Asp Leu
 3440 3445 3450
 His Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe Asn
 3455 3460 3465
 Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys
 3470 3475 3480
 Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu
 3485 3490 3495
 Leu Arg Pro Glu Lys His Gly Ala Ala Thr Gly Val Asp Ala Ile
 3500 3505 3510
 Cys Thr Leu Arg Leu Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu
 3515 3520 3525
 Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Ser Val Thr Glu
 3530 3535 3540
 Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly
 3545 3550 3555
 Phe Thr His Arg Ser Ser Val Pro Thr Thr Ser Ile Pro Gly Thr
 3560 3565 3570
 Ser Ala Val His Leu Glu Thr Ser Gly Thr Pro Ala Ser Leu Pro
 3575 3580 3585
 Gly His Thr Ala Pro Gly Pro Leu Leu Val Pro Phe Thr Leu Asn
 3590 3595 3600
 Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro
 3605 3610 3615
 Gly Ser Arg Lys Phe Ser Thr Thr Glu Arg Val Leu Gln Gly Leu
 3620 3625 3630
 Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu Tyr Ser
 3635 3640 3645
 Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala Ala
 3650 3655 3660

Thr Arg Val Asp Ala Val Cys Thr His Arg Pro Asp Pro Lys Ser
 3665 3670 3675
 Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Lys Leu Ser Gln Leu
 3680 3685 3690
 Thr His Gly Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg His
 3695 3700 3705
 Ser Leu Tyr Val Asn Gly Phe Thr His Gln Ser Ser Met Thr Thr
 3710 3715 3720
 Thr Arg Thr Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser Arg
 3725 3730 3735
 Thr Pro Ala Ser Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu
 3740 3745 3750
 Val Leu Phe Thr Ile Asn Phe Thr Ile Thr Asn Gln Arg Tyr Glu
 3755 3760 3765
 Glu Asn Met His His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu
 3770 3775 3780
 Arg Val Leu Gln Gly Leu Leu Arg Pro Val Phe Lys Asn Thr Ser
 3785 3790 3795
 Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro
 3800 3805 3810
 Lys Lys Asp Gly Ala Ala Thr Lys Val Asp Ala Ile Cys Thr Tyr
 3815 3820 3825
 Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr
 3830 3835 3840
 Trp Glu Leu Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro
 3845 3850 3855
 Tyr Thr Gln Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His
 3860 3865 3870
 Arg Ser Ser Val Pro Thr Thr Ser Ile Pro Gly Thr Ser Ala Val
 3875 3880 3885
 His Leu Glu Thr Ser Gly Thr Pro Ala Ser Leu Pro Gly His Thr
 3890 3895 3900
 Ala Pro Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile
 3905 3910 3915
 Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro Gly Ser Arg
 3920 3925 3930
 Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Lys Pro
 3935 3940 3945
 Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg

3950	3955	3960
Leu Thr 3965	Leu Leu Arg Pro Glu 3970	Lys Arg Gly Ala Ala Thr Gly Val 3975
Asp Thr 3980	Ile Cys Thr His Arg 3985	Leu Asp Pro Leu Asn Pro Gly Leu 3990
Asp Arg 3995	Glu Gln Leu Tyr Trp 4000	Glu Leu Ser Lys Leu Thr Arg Gly 4005
Ile Ile 4010	Glu Leu Gly Pro Tyr 4015	Leu Leu Asp Arg Gly Ser Leu Tyr 4020
Val Asn 4025	Gly Phe Thr His Arg 4030	Thr Ser Val Pro Thr Thr Ser Thr 4035
Pro Gly 4040	Thr Ser Thr Val Asp 4045	Leu Gly Thr Ser Gly Thr Pro Phe 4050
Ser Leu 4055	Pro Ser Pro Ala Xaa 4060	Xaa Xaa Pro Leu Leu Xaa Pro Phe 4065
Thr Leu 4070	Asn Phe Thr Ile Thr 4075	Asn Leu Xaa Tyr Glu Glu Xaa Met 4080
Xaa Xaa 4085	Pro Gly Ser Arg Lys 4090	Phe Asn Thr Thr Glu Arg Val Leu 4095
Gln Thr 4100	Leu Leu Gly Pro Met 4105	Phe Lys Asn Thr Ser Val Gly Leu 4110
Leu Tyr 4115	Ser Gly Cys Arg Leu 4120	Thr Leu Leu Arg Ser Glu Lys Asp 4125
Gly Ala 4130	Ala Thr Gly Val Asp 4135	Ala Ile Cys Thr His Arg Leu Asp 4140
Pro Lys 4145	Ser Pro Gly Val Asp 4150	Arg Glu Gln Leu Tyr Trp Glu Leu 4155
Ser Gln 4160	Leu Thr Asn Gly Ile 4165	Lys Glu Leu Gly Pro Tyr Thr Leu 4170
Asp Arg 4175	Asn Ser Leu Tyr Val 4180	Asn Gly Phe Thr His Trp Ile Pro 4185
Val Pro 4190	Thr Ser Ser Thr Pro 4195	Gly Thr Ser Thr Val Asp Leu Gly 4200
Ser Gly 4205	Thr Pro Ser Leu Pro 4210	Ser Ser Pro Thr Thr Ala Gly Pro 4215
Leu Leu 4220	Val Pro Phe Thr Leu 4225	Asn Phe Thr Ile Thr Asn Leu Lys 4230
Tyr Glu 4235	Glu Asp Met His Cys 4240	Pro Gly Ser Arg Lys Phe Asn Thr 4245

Thr Glu	Arg Val	Leu Gln	Ser	Leu Leu	Gly Pro	Met	Phe Lys	Asn	
4250			4255			4260			
Thr Ser	Val Gly	Pro Leu	Tyr	Ser Gly	Cys Arg	Leu	Thr Leu	Leu	
4265			4270			4275			
Arg Ser	Glu Lys	Asp Gly	Ala	Ala Thr	Gly Val	Asp	Ala Ile	Cys	
4280			4285			4290			
Thr His	Arg Leu	Asp Pro	Lys	Ser Pro	Gly Val	Asp	Arg Glu	Gln	
4295			4300			4305			
Leu Tyr	Trp Glu	Leu Ser	Gln	Leu Thr	Asn Gly	Ile	Lys Glu	Leu	
4310			4315			4320			
Gly Pro	Tyr Thr	Leu Asp	Arg	Asn Ser	Leu Tyr	Val	Asn Gly	Phe	
4325			4330			4335			
Thr His	Gln Thr	Ser Ala	Pro	Asn Thr	Ser Thr	Pro	Gly Thr	Ser	
4340			4345			4350			
Thr Val	Asp Leu	Gly Thr	Ser	Gly Thr	Pro Ser	Ser	Leu Pro	Ser	
4355			4360			4365			
Pro Thr	Xaa Xaa	Xaa Pro	Leu	Leu Xaa	Pro Phe	Thr	Leu Asn	Phe	
4370			4375			4380			
Thr Ile	Thr Asn	Leu Xaa	Tyr	Glu Glu	Xaa Met	Xaa	Xaa Pro	Gly	
4385			4390			4395			
Ser Arg	Lys Phe	Asn Thr	Thr	Glu Arg	Val Leu	Gln	Gly Leu	Leu	
4400			4405			4410			
Xaa Pro	Xaa Phe	Lys Xaa	Thr	Ser Val	Gly Xaa	Leu	Tyr Ser	Gly	
4415			4420			4425			
Cys Arg	Leu Thr	Leu Leu	Arg	Xaa Glu	Lys Xaa	Xaa	Ala Ala	Thr	
4430			4435			4440			
Xaa Val	Asp Xaa	Xaa Cys	Xaa	Xaa Xaa	Xaa Asp	Pro	Xaa Xaa	Pro	
4445			4450			4455			
Gly Leu	Asp Arg	Glu Xaa	Leu	Tyr Trp	Glu Leu	Ser	Xaa Leu	Thr	
4460			4465			4470			
Xaa Xaa	Ile Xaa	Glu Leu	Gly	Pro Tyr	Xaa Leu	Asp	Arg Xaa	Ser	
4475			4480			4485			
Leu Tyr	Val Asn	Gly Phe	Thr	His Trp	Ile Pro	Val	Pro Thr	Ser	
4490			4495			4500			
Ser Thr	Pro Gly	Thr Ser	Thr	Val Asp	Leu Gly	Ser	Gly Thr	Pro	
4505			4510			4515			
Ser Ser	Leu Pro	Ser Pro	Thr	Thr Ala	Gly Pro	Leu	Leu Val	Pro	
4520			4525			4530			
Phe Thr	Leu Asn	Phe Thr	Ile	Thr Asn	Leu Lys	Tyr	Glu Glu	Asp	
4535			4540			4545			

Met	His	Cys	Pro	Gly	Ser	Arg	Lys	Phe	Asn	Thr	Thr	Glu	Arg	Val
4550						4555					4560			
Leu	Gln	Ser	Leu	Leu	Gly	Pro	Met	Phe	Lys	Asn	Thr	Ser	Val	Gly
4565						4570					4575			
Pro	Leu	Tyr	Ser	Gly	Cys	Arg	Leu	Thr	Ser	Leu	Arg	Ser	Glu	Lys
4580						4585					4590			
Asp	Gly	Ala	Ala	Thr	Gly	Val	Asp	Ala	Ile	Cys	Thr	His	Arg	Val
4595						4600					4605			
Asp	Pro	Lys	Ser	Pro	Gly	Val	Asp	Arg	Glu	Gln	Leu	Tyr	Trp	Glu
4610						4615					4620			
Leu	Ser	Gln	Leu	Thr	Asn	Gly	Ile	Lys	Glu	Leu	Gly	Pro	Tyr	Thr
4625						4630					4635			
Leu	Asp	Arg	Asn	Ser	Leu	Tyr	Val	Asn	Gly	Phe	Thr	His	Gln	Thr
4640						4645					4650			
Ser	Ala	Pro	Asn	Thr	Ser	Thr	Pro	Gly	Thr	Ser	Thr	Val	Asp	Leu
4655						4660					4665			
Gly	Thr	Ser	Gly	Thr	Pro	Ser	Ser	Leu	Pro	Ser	Pro	Thr	Ser	Ala
4670						4675					4680			
Gly	Pro	Leu	Leu	Val	Pro	Phe	Thr	Leu	Asn	Phe	Thr	Ile	Thr	Asn
4685						4690					4695			
Leu	Gln	Tyr	Glu	Glu	Asp	Met	His	His	Pro	Gly	Ser	Arg	Lys	Phe
4700						4705					4710			
Asn	Thr	Thr	Glu	Arg	Val	Leu	Gln	Gly	Leu	Leu	Gly	Pro	Met	Phe
4715						4720					4725			
Lys	Asn	Thr	Ser	Val	Gly	Leu	Leu	Tyr	Ser	Gly	Cys	Arg	Leu	Thr
4730						4735					4740			
Leu	Leu	Arg	Pro	Glu	Lys	Asn	Gly	Ala	Ala	Thr	Gly	Met	Asp	Ala
4745						4750					4755			
Ile	Cys	Thr	His	Arg	Leu	Asp	Pro	Lys	Ser	Pro	Gly	Leu	Asp	Arg
4760						4765					4770			
Glu	Xaa	Leu	Tyr	Trp	Glu	Leu	Ser	Xaa	Leu	Thr	Xaa	Xaa	Ile	Xaa
4775						4780					4785			
Glu	Leu	Gly	Pro	Tyr	Xaa	Leu	Asp	Arg	Xaa	Ser	Leu	Tyr	Val	Asn
4790						4795					4800			
Gly	Phe	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Thr	Ser	Thr	Pro	Gly
4805						4810					4815			
Thr	Ser	Xaa	Val	Xaa	Leu	Xaa	Thr	Ser	Gly	Thr	Pro	Xaa	Xaa	Xaa
4820						4825					4830			
Pro	Xaa	Xaa	Thr	Xaa	Xaa	Xaa	Pro	Leu	Leu	Xaa	Pro	Phe	Thr	Leu

4835	4840	4845
Asn Phe Thr Ile Thr Asn	Leu Xaa Tyr Glu Glu Xaa	Met Xaa Xaa
4850	4855	4860
Pro Gly Ser Arg Lys Phe	Asn Thr Thr Glu Arg Val	Leu Gln Gly
4865	4870	4875
Leu Leu Lys Pro Leu Phe	Arg Asn Ser Ser Leu Glu	Tyr Leu Tyr
4880	4885	4890
Ser Gly Cys Arg Leu Ala	Ser Leu Arg Pro Glu Lys	Asp Ser Ser
4895	4900	4905
Ala Met Ala Val Asp Ala	Ile Cys Thr His Arg Pro	Asp Pro Glu
4910	4915	4920
Asp Leu Gly Leu Asp Arg	Glu Arg Leu Tyr Trp Glu	Leu Ser Asn
4925	4930	4935
Leu Thr Asn Gly Ile Gln	Glu Leu Gly Pro Tyr Thr	Leu Asp Arg
4940	4945	4950
Asn Ser Leu Tyr Val Asn	Gly Phe Thr His Arg Ser	Ser Met Pro
4955	4960	4965
Thr Thr Ser Thr Pro Gly	Thr Ser Thr Val Asp Val	Gly Thr Ser
4970	4975	4980
Gly Thr Pro Ser Ser Ser	Pro Ser Pro Thr Thr Ala	Gly Pro Leu
4985	4990	4995
Leu Ile Pro Phe Thr Leu	Asn Phe Thr Ile Thr Asn	Leu Gln Tyr
5000	5005	5010
Gly Glu Asp Met Gly His	Pro Gly Ser Arg Lys Phe	Asn Thr Thr
5015	5020	5025
Glu Arg Val Leu Gln Gly	Leu Leu Gly Pro Ile Phe	Lys Asn Thr
5030	5035	5040
Ser Val Gly Pro Leu Tyr	Ser Gly Cys Arg Leu Thr	Ser Leu Arg
5045	5050	5055
Ser Glu Lys Asp Gly Ala	Ala Thr Gly Val Asp Ala	Ile Cys Ile
5060	5065	5070
His His Leu Asp Pro Lys	Ser Pro Gly Leu Asn Arg	Glu Arg Leu
5075	5080	5085
Tyr Trp Glu Leu Ser Gln	Leu Thr Asn Gly Ile Lys	Glu Leu Gly
5090	5095	5100
Pro Tyr Thr Leu Asp Arg	Asn Ser Leu Tyr Val Asn	Gly Phe Thr
5105	5110	5115
His Arg Thr Ser Val Pro	Thr Thr Ser Thr Pro Gly	Thr Ser Thr
5120	5125	5130

Val Asp 5135	Leu Gly Thr Ser 5140	Gly Thr Pro Phe Ser 5145	Leu Pro Ser Pro
Ala Thr 5150	Ala Gly Pro Leu 5155	Leu Val Leu Phe Thr 5160	Leu Asn Phe Thr
Ile Thr 5165	Asn Leu Lys Tyr 5170	Glu Glu Asp Met His 5175	Arg Pro Gly Ser
Arg Lys 5180	Phe Asn Thr Thr 5185	Glu Arg Val Leu Gln 5190	Thr Leu Leu Gly
Pro Met 5195	Phe Lys Asn Thr 5200	Ser Val Gly Leu Leu 5205	Tyr Ser Gly Cys
Arg Leu 5210	Thr Leu Leu Arg 5215	Glu Lys Asp Gly 5220	Ala Ala Thr Gly
Val Asp 5225	Ala Ile Cys Thr 5230	His Arg Leu Asp Pro 5235	Lys Ser Pro Gly
Leu Asp 5240	Arg Glu Xaa Leu 5245	Tyr Trp Glu Leu Ser 5250	Xaa Leu Thr Xaa
Xaa Ile 5255	Xaa Glu Leu Gly 5260	Pro Tyr Xaa Leu Asp 5265	Arg Xaa Ser Leu
Tyr Val 5270	Asn Gly Phe Xaa 5275	Xaa Xaa Xaa Xaa Xaa 5280	Xaa Thr Ser
Thr Pro 5285	Gly Thr Ser Xaa 5290	Val Xaa Leu Xaa Thr 5295	Ser Gly Thr Pro
Xaa Xaa 5300	Xaa Pro Xaa Xaa 5305	Thr Xaa Xaa Xaa Pro 5310	Leu Leu Xaa Pro
Phe Thr 5315	Leu Asn Phe Thr 5320	Ile Thr Asn Leu Xaa 5325	Tyr Glu Glu Xaa
Met Xaa 5330	Xaa Pro Gly Ser 5335	Arg Lys Phe Asn Thr 5340	Thr Glu Arg Val
Leu Gln 5345	Gly Leu Leu Arg 5350	Pro Val Phe Lys Asn 5355	Thr Ser Val Gly
Pro Leu 5360	Tyr Ser Gly Cys 5365	Arg Leu Thr Leu Leu 5370	Arg Pro Lys Lys
Asp Gly 5375	Ala Ala Thr Lys 5380	Val Asp Ala Ile Cys 5385	Thr Tyr Arg Pro
Asp Pro 5390	Lys Ser Pro Gly 5395	Leu Asp Arg Glu Gln 5400	Leu Tyr Trp Glu
Leu Ser 5405	Gln Leu Thr His 5410	Ser Ile Thr Glu Leu 5415	Gly Pro Tyr Thr
Gln Asp 5420	Arg Asp Ser Leu 5425	Tyr Val Asn Gly Phe 5430	Thr His Arg Ser

Ser Val 5435	Pro Thr Thr Ser 5440	Ile 5440	Pro Gly Thr Ser 5445	Ala 5445	Val His Leu
Glu Thr 5450	Thr Gly Thr Pro 5455	Ser 5455	Ser Phe Pro Gly 5460	His 5460	Thr Glu Pro
Gly Pro 5465	Leu Leu Ile Pro 5470	Phe 5470	Thr Phe Asn Phe 5475	Thr 5475	Ile Thr Asn
Leu Arg 5480	Tyr Glu Glu Asn 5485	Met 5485	Gln His Pro Gly 5490	Ser 5490	Arg Lys Phe
Asn Thr 5495	Thr Glu Arg Val 5500	Leu 5500	Gln Gly Leu Leu 5505	Thr 5505	Pro Leu Phe
Lys Asn 5510	Thr Ser Val Gly 5515	Pro 5515	Leu Tyr Ser Gly 5520	Cys 5520	Arg Leu Thr
Leu Leu 5525	Arg Pro Glu Lys 5530	Gln 5530	Glu Ala Ala Thr 5535	Gly 5535	Val Asp Thr
Ile Cys 5540	Thr His Arg Val 5545	Asp 5545	Pro Ile Gly Pro 5550	Gly 5550	Leu Asp Arg
Glu Arg 5555	Leu Tyr Trp Glu 5560	Leu 5560	Ser Gln Leu Thr 5565	Asn 5565	Ser Ile Thr
Glu Leu 5570	Gly Pro Tyr Thr 5575	Leu 5575	Asp Arg Asp Ser 5580	Leu 5580	Tyr Val Asp
Gly Phe 5585	Asn Pro Trp Ser 5590	Ser 5590	Val Pro Thr Thr 5595	Ser 5595	Thr Pro Gly
Thr Ser 5600	Thr Val His Leu 5605	Ala 5605	Thr Ser Gly Thr 5610	Pro 5610	Ser Pro Leu
Pro Gly 5615	His Thr Ala Pro 5620	Val 5620	Pro Leu Leu Ile 5625	Pro 5625	Phe Thr Leu
Asn Phe 5630	Thr Ile Thr Asp 5635	Leu 5635	His Tyr Glu Glu 5640	Asn 5640	Met Gln His
Pro Gly 5645	Ser Arg Lys Phe 5650	Asn 5650	Thr Thr Glu Arg 5655	Val 5655	Leu Gln Gly
Leu Leu 5660	Lys Pro Leu Phe 5665	Lys 5665	Ser Thr Ser Val 5670	Gly 5670	Pro Leu Tyr
Ser Gly 5675	Cys Arg Leu Thr 5680	Leu 5680	Leu Arg Pro Glu 5685	Lys 5685	His Gly Ala
Ala Thr 5690	Gly Val Asp Ala 5695	Ile 5695	Cys Thr Leu Arg 5700	Leu 5700	Asp Pro Thr
Gly Pro 5705	Gly Leu Asp Arg 5710	Glu 5710	Arg Leu Tyr Trp 5715	Glu 5715	Leu Ser Gln
Leu Thr	Asn Ser Ile Thr 5715	Glu	Leu Gly Pro Tyr 5720	Thr	Leu Asp Arg

5720	5725	5730
Asp Ser Leu Tyr Val Asn Gly Phe Asn Pro Trp Ser Ser Val Pro		
5735	5740	5745
Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala Thr Ser		
5750	5755	5760
Gly Thr Pro Ser Ser Leu Pro Gly His Thr Thr Ala Gly Pro Leu		
5765	5770	5775
Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Lys Tyr		
5780	5785	5790
Glu Glu Asp Met His Cys Pro Gly Ser Arg Lys Phe Asn Thr Thr		
5795	5800	5805
Glu Arg Val Leu Gln Ser Leu His Gly Pro Met Phe Lys Asn Thr		
5810	5815	5820
Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg		
5825	5830	5835
Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr		
5840	5845	5850
His Arg Leu Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Xaa Leu		
5855	5860	5865
Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa Glu Leu Gly		
5870	5875	5880
Pro Tyr Xaa Leu Asp Arg Xaa Ser Leu Tyr Val Asn Gly Phe Xaa		
5885	5890	5895
Xaa Xaa Xaa Xaa Xaa Xaa Thr Ser Thr Pro Gly Thr Ser Xaa		
5900	5905	5910
Val Xaa Leu Xaa Thr Ser Gly Thr Pro Xaa Xaa Xaa Pro Xaa Xaa		
5915	5920	5925
Thr Xaa Xaa Xaa Pro Leu Leu Xaa Pro Phe Thr Leu Asn Phe Thr		
5930	5935	5940
Ile Thr Asn Leu Xaa Tyr Glu Glu Xaa Met Xaa Xaa Pro Gly Ser		
5945	5950	5955
Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Xaa		
5960	5965	5970
Pro Xaa Phe Lys Xaa Thr Ser Val Gly Xaa Leu Tyr Ser Gly Cys		
5975	5980	5985
Arg Leu Thr Leu Leu Arg Xaa Glu Lys Xaa Xaa Ala Ala Thr Xaa		
5990	5995	6000
Val Asp Xaa Xaa Cys Xaa Xaa Xaa Xaa Asp Pro Xaa Xaa Pro Gly		
6005	6010	6015

Leu Asp Arg Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr Asn
 6020 6025 6030
 Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu
 6035 6040 6045
 Tyr Val Asn Gly Phe Thr His Arg Ser Ser Met Pro Thr Thr Ser
 6050 6055 6060
 Ile Pro Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly Thr Pro
 6065 6070 6075
 Ala Ser Leu Pro Gly His Thr Ala Pro Gly Pro Leu Leu Val Pro
 6080 6085 6090
 Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp
 6095 6100 6105
 Met Arg His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val
 6110 6115 6120
 Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly
 6125 6130 6135
 Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys
 6140 6145 6150
 Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu
 6155 6160 6165
 Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Xaa Leu Tyr Trp Glu
 6170 6175 6180
 Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa
 6185 6190 6195
 Leu Asp Arg Xaa Ser Leu Tyr Val Asn Gly Phe Xaa Xaa Xaa Xaa
 6200 6205 6210
 Xaa Xaa Xaa Xaa Thr Ser Thr Pro Gly Thr Ser Xaa Val Xaa Leu
 6215 6220 6225
 Xaa Thr Ser Gly Thr Pro Xaa Xaa Xaa Pro Xaa Xaa Thr Xaa Xaa
 6230 6235 6240
 Xaa Pro Leu Leu Xaa Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn
 6245 6250 6255
 Leu Xaa Tyr Glu Glu Xaa Met Xaa Xaa Pro Gly Ser Arg Lys Phe
 6260 6265 6270
 Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Xaa Pro Xaa Phe
 6275 6280 6285
 Lys Xaa Thr Ser Val Gly Xaa Leu Tyr Ser Gly Cys Arg Leu Thr
 6290 6295 6300
 Leu Leu Arg Xaa Glu Lys Xaa Xaa Ala Ala Thr Xaa Val Asp Xaa
 6305 6310 6315

Xaa Cys 6320	Xaa Xaa Xaa Xaa	Asp 6325	Pro Xaa Xaa Pro	Gly 6330	Leu Asp Arg
Glu Xaa 6335	Leu Tyr Trp Glu	Leu 6340	Ser Xaa Leu Thr	Xaa 6345	Xaa Ile Xaa
Glu Leu 6350	Gly Pro Tyr Xaa	Leu 6355	Asp Arg Xaa Ser	Leu 6360	Tyr Val Asn
Gly Phe 6365	His Pro Arg Ser	Ser 6370	Val Pro Thr Thr	Ser 6375	Thr Pro Gly
Thr Ser 6380	Thr Val His Leu	Ala 6385	Thr Ser Gly Thr	Pro 6390	Ser Ser Leu
Pro Gly 6395	His Thr Ala Pro	Val 6400	Pro Leu Leu Ile	Pro 6405	Phe Thr Leu
Asn Phe 6410	Thr Ile Thr Asn	Leu 6415	His Tyr Glu Glu	Asn 6420	Met Gln His
Pro Gly 6425	Ser Arg Lys Phe	Asn 6430	Thr Thr Glu Arg	Val 6435	Leu Gln Gly
Leu Leu 6440	Gly Pro Met Phe	Lys 6445	Asn Thr Ser Val	Gly 6450	Leu Leu Tyr
Ser Gly 6455	Cys Arg Leu Thr	Leu 6460	Leu Arg Pro Glu	Lys 6465	Asn Gly Ala
Ala Thr 6470	Gly Met Asp Ala	Ile 6475	Cys Ser His Arg	Leu 6480	Asp Pro Lys
Ser Pro 6485	Gly Leu Asp Arg	Glu 6490	Xaa Leu Tyr Trp	Glu 6495	Leu Ser Xaa
Leu Thr 6500	Xaa Xaa Ile Xaa	Glu 6505	Leu Gly Pro Tyr	Xaa 6510	Leu Asp Arg
Xaa Ser 6515	Leu Tyr Val Asn	Gly 6520	Phe Xaa Xaa Xaa	Xaa 6525	Xaa Xaa Xaa
Xaa Thr 6530	Ser Thr Pro Gly	Thr 6535	Ser Xaa Val Xaa	Leu 6540	Xaa Thr Ser
Gly Thr 6545	Pro Xaa Xaa Xaa	Pro 6550	Xaa Xaa Thr Xaa	Xaa 6555	Xaa Pro Leu
Leu Xaa 6560	Pro Phe Thr Leu	Asn 6565	Phe Thr Ile Thr	Asn 6570	Leu Xaa Tyr
Glu Glu 6575	Xaa Met Xaa Xaa	Pro 6580	Gly Ser Arg Lys	Phe 6585	Asn Thr Thr
Glu Arg 6590	Val Leu Gln Gly	Leu 6595	Leu Xaa Pro Xaa	Phe 6600	Lys Xaa Thr
Ser Val	Gly Xaa Leu Tyr	Ser	Gly Cys Arg Leu Thr	Leu Leu Arg	

6605		6610		6615
Xaa Glu Lys Xaa Xaa Ala	Ala	Thr Xaa Val Asp Xaa	Xaa Cys Xaa	
6620	6625	6630		
Xaa Xaa Xaa Asp Pro Xaa	Xaa	Pro Gly Leu Asp Arg	Glu Xaa Leu	
6635	6640	6645		
Tyr Trp Glu Leu Ser Xaa	Leu	Thr Xaa Xaa Ile Xaa	Glu Leu Gly	
6650	6655	6660		
Pro Tyr Xaa Leu Asp Arg	Xaa	Ser Leu Tyr Val Asn	Gly Phe Thr	
6665	6670	6675		
His Gln Asn Ser Val Pro	Thr	Thr Ser Thr Pro Gly	Thr Ser Thr	
6680	6685	6690		
Val Tyr Trp Ala Thr Thr	Gly	Thr Pro Ser Ser Phe	Pro Gly His	
6695	6700	6705		
Thr Glu Pro Gly Pro Leu	Leu	Ile Pro Phe Thr Phe	Asn Phe Thr	
6710	6715	6720		
Ile Thr Asn Leu His Tyr	Glu	Glu Asn Met Gln His	Pro Gly Ser	
6725	6730	6735		
Arg Lys Phe Asn Thr Thr	Glu	Arg Val Leu Gln Gly	Leu Leu Thr	
6740	6745	6750		
Pro Leu Phe Lys Asn Thr	Ser	Val Gly Pro Leu Tyr	Ser Gly Cys	
6755	6760	6765		
Arg Leu Thr Leu Leu Arg	Pro	Glu Lys Gln Glu Ala	Ala Thr Gly	
6770	6775	6780		
Val Asp Thr Ile Cys Thr	His	Arg Val Asp Pro Ile	Gly Pro Gly	
6785	6790	6795		
Leu Asp Arg Glu Xaa Leu	Tyr	Trp Glu Leu Ser Xaa	Leu Thr Xaa	
6800	6805	6810		
Xaa Ile Xaa Glu Leu Gly	Pro	Tyr Xaa Leu Asp Arg	Xaa Ser Leu	
6815	6820	6825		
Tyr Val Asn Gly Phe Xaa	Xaa	Xaa Xaa Xaa Xaa Xaa	Xaa Thr Ser	
6830	6835	6840		
Thr Pro Gly Thr Ser Xaa	Val	Xaa Leu Xaa Thr Ser	Gly Thr Pro	
6845	6850	6855		
Xaa Xaa Xaa Pro Xaa Xaa	Thr	Xaa Xaa Xaa Pro Leu	Leu Xaa Pro	
6860	6865	6870		
Phe Thr Leu Asn Phe Thr	Ile	Thr Asn Leu Xaa Tyr	Glu Glu Xaa	
6875	6880	6885		
Met Xaa Xaa Pro Gly Ser	Arg	Lys Phe Asn Thr Thr	Glu Arg Val	
6890	6895	6900		

Leu	Gln	Gly	Leu	Leu	Xaa	Pro	Xaa	Phe	Lys	Xaa	Thr	Ser	Val	Gly
6905						6910					6915			
Xaa	Leu	Tyr	Ser	Gly	Cys	Arg	Leu	Thr	Leu	Leu	Arg	Xaa	Glu	Lys
6920						6925					6930			
Xaa	Xaa	Ala	Ala	Thr	Xaa	Val	Asp	Xaa	Xaa	Cys	Xaa	Xaa	Xaa	Xaa
6935						6940					6945			
Asp	Pro	Xaa	Xaa	Pro	Gly	Leu	Asp	Arg	Glu	Xaa	Leu	Tyr	Trp	Glu
6950						6955					6960			
Leu	Ser	Xaa	Leu	Thr	Xaa	Xaa	Ile	Xaa	Glu	Leu	Gly	Pro	Tyr	Xaa
6965						6970					6975			
Leu	Asp	Arg	Xaa	Ser	Leu	Tyr	Val	Asn	Gly	Phe	Thr	His	Arg	Ser
6980						6985					6990			
Ser	Val	Pro	Thr	Thr	Ser	Ser	Pro	Gly	Thr	Ser	Thr	Val	His	Leu
6995						7000					7005			
Ala	Thr	Ser	Gly	Thr	Pro	Ser	Ser	Leu	Pro	Gly	His	Thr	Ala	Pro
7010						7015					7020			
Val	Pro	Leu	Leu	Ile	Pro	Phe	Thr	Leu	Asn	Phe	Thr	Ile	Thr	Asn
7025						7030					7035			
Leu	His	Tyr	Glu	Glu	Asn	Met	Gln	His	Pro	Gly	Ser	Arg	Lys	Phe
7040						7045					7050			
Asn	Thr	Thr	Glu	Arg	Val	Leu	Gln	Gly	Leu	Leu	Lys	Pro	Leu	Phe
7055						7060					7065			
Lys	Ser	Thr	Ser	Val	Gly	Pro	Leu	Tyr	Ser	Gly	Cys	Arg	Leu	Thr
7070						7075					7080			
Leu	Leu	Arg	Pro	Glu	Lys	His	Gly	Ala	Ala	Thr	Gly	Val	Asp	Ala
7085						7090					7095			
Ile	Cys	Thr	Leu	Arg	Leu	Asp	Pro	Thr	Gly	Pro	Gly	Leu	Asp	Arg
7100						7105					7110			
Glu	Xaa	Leu	Tyr	Trp	Glu	Leu	Ser	Xaa	Leu	Thr	Xaa	Xaa	Ile	Xaa
7115						7120					7125			
Glu	Leu	Gly	Pro	Tyr	Xaa	Leu	Asp	Arg	Xaa	Ser	Leu	Tyr	Val	Asn
7130						7135					7140			
Gly	Phe	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Thr	Ser	Thr	Pro	Gly
7145						7150					7155			
Thr	Ser	Xaa	Val	Xaa	Leu	Xaa	Thr	Ser	Gly	Thr	Pro	Xaa	Xaa	Xaa
7160						7165					7170			
Pro	Xaa	Xaa	Thr	Xaa	Xaa	Xaa	Pro	Leu	Leu	Xaa	Pro	Phe	Thr	Leu
7175						7180					7185			
Asn	Phe	Thr	Ile	Thr	Asn	Leu	Xaa	Tyr	Glu	Glu	Xaa	Met	Xaa	Xaa
7190						7195					7200			

Pro Gly	Ser Arg	Lys Phe	Asn Thr	Thr Thr	Glu Arg	Val Leu	Gln Gly		
7205			7210			7215			
Leu Leu	Xaa Pro	Xaa Phe	Lys Xaa	Thr Ser	Val Gly	Xaa Leu	Tyr		
7220			7225			7230			
Ser Gly	Cys Arg	Leu Thr	Leu Leu	Arg Xaa	Glu Lys	Xaa Xaa	Ala		
7235			7240			7245			
Ala Thr	Xaa Val	Asp Xaa	Xaa Cys	Xaa Xaa	Xaa Xaa	Asp Pro	Xaa		
7250			7255			7260			
Xaa Pro	Gly Leu	Asp Arg	Glu Xaa	Leu Tyr	Trp Glu	Leu Ser	Xaa		
7265			7270			7275			
Leu Thr	Xaa Xaa	Ile Xaa	Glu Leu	Gly Pro	Tyr Xaa	Leu Asp	Arg		
7280			7285			7290			
Xaa Ser	Leu Tyr	Val Asn	Gly Phe	Thr His	Arg Thr	Ser Val	Pro		
7295			7300			7305			
Thr Thr	Ser Thr	Pro Gly	Thr Ser	Thr Val	His Leu	Ala Thr	Ser		
7310			7315			7320			
Gly Thr	Pro Ser	Ser Leu	Pro Gly	His Thr	Ala Pro	Val Pro	Leu		
7325			7330			7335			
Leu Ile	Pro Phe	Thr Leu	Asn Phe	Thr Ile	Thr Asn	Leu Gln	Tyr		
7340			7345			7350			
Glu Glu	Asp Met	His Arg	Pro Gly	Ser Arg	Lys Phe	Asn Thr	Thr		
7355			7360			7365			
Glu Arg	Val Leu	Gln Gly	Leu Leu	Ser Pro	Ile Phe	Lys Asn	Ser		
7370			7375			7380			
Ser Val	Gly Pro	Leu Tyr	Ser Gly	Cys Arg	Leu Thr	Ser Leu	Arg		
7385			7390			7395			
Pro Glu	Lys Asp	Gly Ala	Ala Thr	Gly Met	Asp Ala	Val Cys	Leu		
7400			7405			7410			
Tyr His	Pro Asn	Pro Lys	Arg Pro	Gly Leu	Asp Arg	Glu Gln	Leu		
7415			7420			7425			
Tyr Cys	Glu Leu	Ser Gln	Leu Thr	His Asn	Ile Thr	Glu Leu	Gly		
7430			7435			7440			
Pro Tyr	Ser Leu	Asp Arg	Asp Ser	Leu Tyr	Val Asn	Gly Phe	Thr		
7445			7450			7455			
His Gln	Asn Ser	Val Pro	Thr Thr	Ser Thr	Pro Gly	Thr Ser	Thr		
7460			7465			7470			
Val Tyr	Trp Ala	Thr Thr	Gly Thr	Pro Ser	Ser Phe	Pro Gly	His		
7475			7480			7485			
Thr Xaa	Xaa Xaa	Pro Leu	Leu Xaa	Pro Phe	Thr Leu	Asn Phe	Thr		

7490		7495		7500
Ile Thr Asn Leu Xaa Tyr	Glu	Glu Xaa Met Xaa Xaa	Pro Gly Ser	
7505	7510	7515		
Arg Lys Phe Asn Thr Thr	Glu	Arg Val Leu Gln Gly	Leu Leu Xaa	
7520	7525	7530		
Pro Xaa Phe Lys Xaa Thr	Ser	Val Gly Xaa Leu Tyr	Ser Gly Cys	
7535	7540	7545		
Arg Leu Thr Leu Leu Arg	Xaa	Glu Lys Xaa Xaa Ala	Ala Thr Xaa	
7550	7555	7560		
Val Asp Xaa Xaa Cys Xaa	Xaa	Xaa Xaa Asp Pro Xaa	Xaa Pro Gly	
7565	7570	7575		
Leu Asp Arg Glu Xaa Leu	Tyr	Trp Glu Leu Ser Xaa	Leu Thr Xaa	
7580	7585	7590		
Xaa Ile Xaa Glu Leu Gly	Pro	Tyr Xaa Leu Asp Arg	Xaa Ser Leu	
7595	7600	7605		
Tyr Val Asn Gly Phe Thr	His	Trp Ser Ser Gly Leu	Thr Thr Ser	
7610	7615	7620		
Thr Pro Trp Thr Ser Thr	Val	Asp Leu Gly Thr Ser	Gly Thr Pro	
7625	7630	7635		
Ser Pro Val Pro Ser Pro	Thr	Thr Ala Gly Pro Leu	Leu Val Pro	
7640	7645	7650		
Phe Thr Leu Asn Phe Thr	Ile	Thr Asn Leu Gln Tyr	Glu Glu Asp	
7655	7660	7665		
Met His Arg Pro Gly Ser	Arg	Lys Phe Asn Ala Thr	Glu Arg Val	
7670	7675	7680		
Leu Gln Gly Leu Leu Ser	Pro	Ile Phe Lys Asn Thr	Ser Val Gly	
7685	7690	7695		
Pro Leu Tyr Ser Gly Cys	Arg	Leu Thr Leu Leu Arg	Pro Glu Lys	
7700	7705	7710		
Gln Glu Ala Ala Thr Gly	Val	Asp Thr Ile Cys Thr	His Arg Val	
7715	7720	7725		
Asp Pro Ile Gly Pro Gly	Leu	Asp Arg Glu Xaa Leu	Tyr Trp Glu	
7730	7735	7740		
Leu Ser Xaa Leu Thr Xaa	Xaa	Ile Xaa Glu Leu Gly	Pro Tyr Xaa	
7745	7750	7755		
Leu Asp Arg Xaa Ser Leu	Tyr	Val Asn Gly Phe Xaa	Xaa Xaa Xaa	
7760	7765	7770		
Xaa Xaa Xaa Xaa Thr Ser	Thr	Pro Gly Thr Ser Xaa	Val Xaa Leu	
7775	7780	7785		

Xaa Thr	Ser Gly Thr	Pro Xaa	Xaa Xaa Pro Xaa Xaa	Thr Xaa Xaa	7790	7795	7800
Xaa Pro	Leu Leu Xaa	Pro Phe	Thr Leu Asn Phe	Thr Ile Thr Asn	7805	7810	7815
Leu Xaa	Tyr Glu Glu Xaa	Met	Xaa Xaa Pro Gly Ser	Arg Lys Phe	7820	7825	7830
Asn Thr	Thr Glu Arg Val	Leu	Gln Gly Leu Leu Xaa	Pro Xaa Phe	7835	7840	7845
Lys Xaa	Thr Ser Val Gly	Xaa	Leu Tyr Ser Gly Cys	Arg Leu Thr	7850	7855	7860
Leu Leu	Arg Xaa Glu Lys	Xaa	Xaa Ala Ala Thr Xaa	Val Asp Xaa	7865	7870	7875
Xaa Cys	Xaa Xaa Xaa Xaa	Asp	Pro Xaa Xaa Pro Gly	Leu Asp Arg	7880	7885	7890
Glu Xaa	Leu Tyr Trp Glu	Leu	Ser Xaa Leu Thr Xaa	Xaa Ile Xaa	7895	7900	7905
Glu Leu	Gly Pro Tyr Xaa	Leu	Asp Arg Xaa Ser Leu	Tyr Val Asn	7910	7915	7920
Gly Phe	Thr His Arg Ser	Phe	Gly Leu Thr Thr Ser	Thr Pro Trp	7925	7930	7935
Thr Ser	Thr Val Asp Leu	Gly	Thr Ser Gly Thr Pro	Ser Pro Val	7940	7945	7950
Pro Ser	Pro Thr Thr Ala	Gly	Pro Leu Leu Val Pro	Phe Thr Leu	7955	7960	7965
Asn Phe	Thr Ile Thr Asn	Leu	Gln Tyr Glu Glu Asp	Met His Arg	7970	7975	7980
Pro Gly	Ser Arg Lys Phe	Asn	Thr Thr Glu Arg Val	Leu Gln Gly	7985	7990	7995
Leu Leu	Thr Pro Leu Phe	Arg	Asn Thr Ser Val Ser	Ser Leu Tyr	8000	8005	8010
Ser Gly	Cys Arg Leu Thr	Leu	Leu Arg Pro Glu Lys	Asp Gly Ala	8015	8020	8025
Ala Thr	Arg Val Asp Ala	Val	Cys Thr His Arg Pro	Asp Pro Lys	8030	8035	8040
Ser Pro	Gly Leu Asp Arg	Glu	Xaa Leu Tyr Trp Glu	Leu Ser Xaa	8045	8050	8055
Leu Thr	Xaa Xaa Ile Xaa	Glu	Leu Gly Pro Tyr Xaa	Leu Asp Arg	8060	8065	8070
Xaa Ser	Leu Tyr Val Asn	Gly	Phe Xaa Xaa Xaa Xaa	Xaa Xaa Xaa	8075	8080	8085

Xaa Thr 8090	Ser Thr Pro Gly Thr 8095	Ser Xaa Val Xaa Leu 8100	Xaa Thr Ser
Gly Thr 8105	Pro Xaa Xaa Xaa Pro 8110	Xaa Xaa Thr Xaa Xaa 8115	Xaa Pro Leu
Leu Xaa 8120	Pro Phe Thr Leu Asn 8125	Phe Thr Ile Thr Asn 8130	Leu Xaa Tyr
Glu Glu 8135	Xaa Met Xaa Xaa Pro 8140	Gly Ser Arg Lys Phe 8145	Asn Thr Thr
Glu Arg 8150	Val Leu Gln Gly Leu 8155	Leu Xaa Pro Xaa Phe 8160	Lys Xaa Thr
Ser Val 8165	Gly Xaa Leu Tyr Ser 8170	Gly Cys Arg Leu Thr 8175	Leu Leu Arg
Xaa Glu 8180	Lys Xaa Xaa Ala Ala 8185	Thr Xaa Val Asp Xaa 8190	Xaa Cys Xaa
Xaa Xaa 8195	Xaa Asp Pro Xaa Xaa 8200	Pro Gly Leu Asp Arg 8205	Glu Xaa Leu
Tyr Trp 8210	Glu Leu Ser Xaa Leu 8215	Thr Xaa Xaa Ile Xaa 8220	Glu Leu Gly
Pro Tyr 8225	Xaa Leu Asp Arg Xaa 8230	Ser Leu Tyr Val Asn 8235	Gly Phe Thr
His Trp 8240	Ile Pro Val Pro Thr 8245	Ser Ser Thr Pro Gly 8250	Thr Ser Thr
Val Asp 8255	Leu Gly Ser Gly Thr 8260	Pro Ser Ser Leu Pro 8265	Ser Pro Thr
Thr Ala 8270	Gly Pro Leu Leu Val 8275	Pro Phe Thr Leu Asn 8280	Phe Thr Ile
Thr Asn 8285	Leu Gln Tyr Gly Glu 8290	Asp Met Gly His Pro 8295	Gly Ser Arg
Lys Phe 8300	Asn Thr Thr Glu Arg 8305	Val Leu Gln Gly Leu 8310	Leu Gly Pro
Ile Phe 8315	Lys Asn Thr Ser Val 8320	Gly Pro Leu Tyr Ser 8325	Gly Cys Arg
Leu Thr 8330	Ser Leu Arg Ser Glu 8335	Lys Asp Gly Ala Ala 8340	Thr Gly Val
Asp Ala 8345	Ile Cys Ile His His 8350	Leu Asp Pro Lys Ser 8355	Pro Gly Leu
Asp Arg 8360	Glu Xaa Leu Tyr Trp 8365	Glu Leu Ser Xaa Leu 8370	Thr Xaa Xaa
Ile Xaa	Glu Leu Gly Pro Tyr	Xaa Leu Asp Arg Xaa	Ser Leu Tyr

8375	8380	8385
Val Asn Gly Phe Xaa Xaa Xaa Xaa Xaa Xaa Xaa Thr Ser Thr		
8390	8395	8400
Pro Gly Thr Ser Xaa Val Xaa Leu Xaa Thr Ser Gly Thr Pro Xaa		
8405	8410	8415
Xaa Xaa Pro Xaa Xaa Thr Xaa Xaa Xaa Pro Leu Leu Xaa Pro Phe		
8420	8425	8430
Thr Leu Asn Phe Thr Ile Thr Asn Leu Xaa Tyr Glu Glu Xaa Met		
8435	8440	8445
Xaa Xaa Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu		
8450	8455	8460
Gln Gly Leu Leu Xaa Pro Xaa Phe Lys Xaa Thr Ser Val Gly Xaa		
8465	8470	8475
Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Xaa Glu Lys Xaa		
8480	8485	8490
Xaa Ala Ala Thr Xaa Val Asp Xaa Xaa Cys Xaa Xaa Xaa Xaa Asp		
8495	8500	8505
Pro Xaa Xaa Pro Gly Leu Asp Arg Glu Xaa Leu Tyr Trp Glu Leu		
8510	8515	8520
Ser Xaa Leu Thr Xaa Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa Leu		
8525	8530	8535
Asp Arg Xaa Ser Leu Tyr Val Asn Gly Phe Thr His Gln Thr Phe		
8540	8545	8550
Ala Pro Asn Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly		
8555	8560	8565
Thr Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr Ser Ala Gly		
8570	8575	8580
Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu		
8585	8590	8595
Gln Tyr Glu Glu Asp Met His His Pro Gly Ser Arg Lys Phe Asn		
8600	8605	8610
Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Met Phe Lys		
8615	8620	8625
Asn Thr Ser Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu		
8630	8635	8640
Leu Arg Pro Glu Lys Asn Gly Ala Ala Thr Arg Val Asp Ala Val		
8645	8650	8655
Cys Thr His Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu		
8660	8665	8670

Xaa Leu	Tyr Trp	Glu Leu	Ser	Xaa Leu	Thr Xaa	Xaa	Ile Xaa	Glu
8675			8680			8685		
Leu Gly	Pro Tyr	Xaa Leu	Asp	Arg Xaa	Ser Leu	Tyr	Val Asn	Gly
8690			8695			8700		
Phe Xaa	Xaa Xaa	Xaa Xaa	Xaa	Xaa Xaa	Thr Ser	Thr	Pro Gly	Thr
8705			8710			8715		
Ser Xaa	Val Xaa	Leu Xaa	Thr	Ser Gly	Thr Pro	Xaa	Xaa Xaa	Pro
8720			8725			8730		
Xaa Xaa	Thr Xaa	Xaa Xaa	Pro	Leu Leu	Xaa Pro	Phe	Thr Leu	Asn
8735			8740			8745		
Phe Thr	Ile Thr	Asn Leu	Xaa	Tyr Glu	Glu Xaa	Met	Xaa Xaa	Pro
8750			8755			8760		
Gly Ser	Arg Lys	Phe Asn	Thr	Thr Glu	Arg Val	Leu	Gln Gly	Leu
8765			8770			8775		
Leu Lys	Pro Leu	Phe Lys	Ser	Thr Ser	Val Gly	Pro	Leu Tyr	Ser
8780			8785			8790		
Gly Cys	Arg Leu	Thr Leu	Leu	Arg Pro	Glu Lys	Asp	Gly Val	Ala
8795			8800			8805		
Thr Arg	Val Asp	Ala Ile	Cys	Thr His	Arg Pro	Asp	Pro Lys	Ile
8810			8815			8820		
Pro Gly	Leu Asp	Arg Gln	Gln	Leu Tyr	Trp Glu	Leu	Ser Gln	Leu
8825			8830			8835		
Thr His	Ser Ile	Thr Glu	Leu	Gly Pro	Tyr Thr	Leu	Asp Arg	Asp
8840			8845			8850		
Ser Leu	Tyr Val	Asn Gly	Phe	Thr Gln	Arg Ser	Ser	Val Pro	Thr
8855			8860			8865		
Thr Ser	Thr Pro	Gly Thr	Phe	Thr Val	Gln Pro	Glu	Thr Ser	Glu
8870			8875			8880		
Thr Pro	Ser Ser	Leu Pro	Gly	Pro Thr	Ala Thr	Gly	Pro Val	Leu
8885			8890			8895		
Leu Pro	Phe Thr	Leu Asn	Phe	Thr Ile	Thr Asn	Leu	Gln Tyr	Glu
8900			8905			8910		
Glu Asp	Met His	Arg Pro	Gly	Ser Arg	Lys Phe	Asn	Thr Thr	Glu
8915			8920			8925		
Arg Val	Leu Gln	Gly Leu	Leu	Met Pro	Leu Phe	Lys	Asn Thr	Ser
8930			8935			8940		
Val Ser	Ser Leu	Tyr Ser	Gly	Cys Arg	Leu Thr	Leu	Leu Arg	Pro
8945			8950			8955		
Glu Lys	Asp Gly	Ala Ala	Thr	Arg Val	Asp Ala	Val	Cys Thr	His
8960			8965			8970		

Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Arg Leu Tyr
 8975 8980 8985
 Trp Lys Leu Ser Gln Leu Thr His Gly Ile Thr Glu Leu Gly Pro
 8990 8995 9000
 Tyr Thr Leu Asp Arg His Ser Leu Tyr Val Asn Gly Phe Thr His
 9005 9010 9015
 Gln Ser Ser Met Thr Thr Thr Arg Thr Pro Asp Thr Ser Thr Met
 9020 9025 9030
 His Leu Ala Thr Ser Arg Thr Pro Ala Ser Leu Ser Gly Pro Thr
 9035 9040 9045
 Thr Ala Ser Pro Leu Leu Val Leu Phe Thr Ile Asn Phe Thr Ile
 9050 9055 9060
 Thr Asn Leu Arg Tyr Glu Glu Asn Met His His Pro Gly Ser Arg
 9065 9070 9075
 Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Arg Pro
 9080 9085 9090
 Val Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg
 9095 9100 9105
 Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala Ala Thr Lys Val
 9110 9115 9120
 Asp Ala Ile Cys Thr Tyr Arg Pro Asp Pro Lys Ser Pro Gly Leu
 9125 9130 9135
 Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His Ser
 9140 9145 9150
 Ile Thr Glu Leu Gly Pro Tyr Thr Gln Asp Arg Asp Ser Leu Tyr
 9155 9160 9165
 Asn Val Gly Phe Thr Gln Arg Ser Ser Val Pro Thr Thr Ser Val
 9170 9175 9180
 Pro Gly Thr Pro Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Val
 9185 9190 9195
 Ser Lys Pro Gly Pro Ser Ala Ala Ser Pro Leu Leu Val Leu Phe
 9200 9205 9210
 Thr Leu Asn Gly Thr Ile Thr Asn Leu Arg Tyr Glu Glu Asn Met
 9215 9220 9225
 Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu
 9230 9235 9240
 Gln Gly Leu Leu Arg Ser Leu Phe Lys Ser Thr Ser Val Gly Pro
 9245 9250 9255
 Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp

9260	9265	9270
Gly Thr Ala Thr Gly Val Asp 9275	Ala Ile Cys Thr His 9280	His Pro Asp 9285
Pro Lys Ser Pro Arg Leu Asp 9290	Arg Glu Gln Leu Tyr 9295	Trp Glu Leu 9300
Ser Gln Leu Thr His Asn Ile 9305	Thr Glu Leu Gly 9310	His Tyr Ala Leu 9315
Asp Asn Asp Ser Leu Phe Val 9320	Asn Gly Phe Thr 9325	His Arg Ser Ser 9330
Val Ser Thr Thr Ser Thr 9335	Pro Gly Thr Pro Thr 9340	Val Tyr Leu Gly 9345
Ala Ser Lys Thr Pro Ala Ser 9350	Ile Phe Gly Pro 9355	Ser Ala Ala Ser 9360
His Leu Leu Ile Leu Phe Thr 9365	Leu Asn Phe Thr 9370	Ile Thr Asn Leu 9375
Arg Tyr Glu Glu Asn Met Trp 9380	Pro Gly Ser Arg 9385	Lys Phe Asn Thr 9390
Thr Glu Arg Val Leu Gln Gly 9395	Leu Leu Arg Pro 9400	Leu Phe Lys Asn 9405
Thr Ser Val Gly Pro Leu Tyr 9410	Ser Gly Ser Arg 9415	Leu Thr Leu Leu 9420
Arg Pro Glu Lys Asp Gly 9425	Glu Ala Thr Gly Val 9430	Asp Ala Ile Cys 9435
Thr His Arg Pro Asp Pro 9440	Thr Gly Pro Gly Leu 9445	Asp Arg Glu Gln 9450
Leu Tyr Leu Glu Leu Ser 9455	Gln Leu Thr His Ser 9460	Ile Thr Glu Leu 9465
Gly Pro Tyr Thr Leu Asp 9470	Arg Asp Ser Leu Tyr 9475	Val Asn Gly Phe 9480
Thr His Arg Ser Ser Val 9485	Pro Thr Thr Ser Thr 9490	Gly Val Val Ser 9495
Glu Glu Pro Phe Thr Leu 9500	Asn Phe Thr Ile Asn 9505	Asn Leu Arg Tyr 9510
Met Ala Asp Met Gly Gln 9515	Pro Gly Ser Leu Lys 9520	Phe Asn Ile Thr 9525
Asp Asn Val Met Lys His 9530	Leu Leu Ser Pro Leu 9535	Phe Gln Arg Ser 9540
Ser Leu Gly Ala Arg Tyr 9545	Thr Gly Cys Arg Val 9550	Ile Ala Leu Arg 9555

Ser Val Lys Asn Gly Ala Glu Thr Arg Val Asp Leu Leu Cys Thr
 9560 9565
 Tyr Leu Gln Pro Leu Ser Gly Pro Gly Leu Pro Ile Lys Gln Val
 9575 9580
 Phe His Glu Leu Ser Gln Gln Thr His Gly Ile Thr Arg Leu Gly
 9590 9595
 Pro Tyr Ser Leu Asp Lys Asp Ser Leu Tyr Leu Asn Gly Tyr Asn
 9605 9610
 Glu Pro Gly Leu Asp Glu Pro Pro Thr Thr Pro Lys Pro Ala Thr
 9620 9625
 Thr Phe Leu Pro Pro Leu Ser Glu Ala Thr Thr Ala Met Gly Tyr
 9635 9640
 His Leu Lys Thr Leu Thr Leu Asn Phe Thr Ile Ser Asn Leu Gln
 9650 9655
 Tyr Ser Pro Asp Met Gly Lys Gly Ser Ala Thr Phe Asn Ser Thr
 9665 9670
 Glu Gly Val Leu Gln His Leu Leu Arg Pro Leu Phe Gln Lys Ser
 9680 9685
 Ser Met Gly Pro Phe Tyr Leu Gly Cys Gln Leu Ile Ser Leu Arg
 9695 9700
 Pro Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Thr Thr Cys Thr
 9710 9715
 Tyr His Pro Asp Pro Val Gly Pro Gly Leu Asp Ile Gln Gln Leu
 9725 9730
 Tyr Trp Glu Leu Ser Gln Leu Thr His Gly Val Thr Gln Leu Gly
 9740 9745
 Phe Tyr Val Leu Asp Arg Asp Ser Leu Phe Ile Asn Gly Tyr Ala
 9755 9760
 Pro Gln Asn Leu Ser Ile Arg Gly Glu Tyr Gln Ile Asn Phe His
 9770 9775
 Ile Val Asn Trp Asn Leu Ser Asn Pro Asp Pro Thr Ser Ser Glu
 9785 9790

Tyr

<210> 147

<211> 1422

<212> DNA

<213> Homo sapiens

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<400> 147
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ctgctcagac ccttgttcca gaagagcagc atgggcccct tctacttggg ttgccaactg      180
atctccctca ggctgagaa ggatggggca gccactggtg tggacaccac ctgcacctac      240
caccctgacc ctgtgggccc cgggctggac atacagcagc tttactggga gctgagtcag      300
ctgaccocatg gtgtcaccca actgggcttc tatgtcctgg acaggatag cctcttcac      360
aatggtcatg cccccagaa tttatcaatc cgggcgaggt accagataaa tttccacatt      420
gtcaactgga acctcagtaa tccagacccc acatcctcag agtacatcac cctgtgagg      480
gacatccagg acaaggtcac cacactctac aaaggcagtc aactacatga cacattccgc      540
ttctgcctgg tcaccaactt gacgatggac tccgtgttgg tcaactgtcaa ggcattgttc      600
tctccaatt tggacccag cctggtggag caagtcttcc tagataagac cctgaatgcc      660
tcattccatt ggctgggctc cacctaccag ttggtggaca tccatgtgac agaaatggag      720
tcactagttt atcaaccaac aagcagctcc agcaccacag acttctacct gaatttcacc      780
atcaccaccc taccatattc ccaggacaaa gccagccag gcaccaccaa ttaccagagg      840
aacaaaagga atattgagga tgcgctcaac caactcttcc gaacacagcag catcaagagt      900
tatttttctg actgtcaagt ttcaacattc aggtctgtcc ccaacaggca ccacaccggg      960
gtggactccc tgtgtaactt ctgccactg gctcggagag tagacagagt tgccatctat      1020
gaggaatttc tgcggatgac ccggaatggt acccagctgc agaacttcac cctggacagg      1080
agcagtgctc ttgtggatgg gtattctccc aacagaaatg agcccttaac tgggaattct      1140
gaccttccct tctgggtgt catctctatc ggcttggcag gactcctggg actcatcaca      1200
tgctgatct gcggtgtcct ggtgaccacc gcgcggcgga agaaggaagg agaatacaac      1260
gtccagcaac agtgcccagg ctactaccag tcacacctag acctggagga tctgcaatga      1320
ctggaaactg ccggtgcctg ggggtgcctt ccccagcca ggtccaaag aagcttggct      1380
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<211> 439

<212> PRT

<400> 148

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35 40 45

Ser Ser Met Gly Pro Phe Tyr Leu Gly Cys Gln Leu Ile Ser Leu Arg
50 55 60

Pro Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Thr Thr Cys Thr Tyr
65 70 75 80

His Pro Asp Pro Val Gly Pro Gly Leu Asp Ile Gln Gln Leu Tyr Trp
85 90 95

Glu Leu Ser Gln Leu Thr His Gly Val Thr Gln Leu Gly Phe Tyr Val
100 105 110

Leu Asp Arg Asp Ser Leu Phe Ile Asn Gly Tyr Ala Pro Gln Asn Leu
115 120 125

Ser Ile Arg Gly Glu Tyr Gln Ile Asn Phe His Ile Val Asn Trp Asn
130 135 140

Leu Ser Asn Pro Asp Pro Thr Ser Ser Glu Tyr Ile Thr Leu Leu Arg
145 150 155 160

Asp Ile Gln Asp Lys Val Thr Thr Leu Tyr Lys Gly Ser Gln Leu His
165 170 175

Asp Thr Phe Arg Phe Cys Leu Val Thr Asn Leu Thr Met Asp Ser Val
180 185 190

Leu Val Thr Val Lys Ala Leu Phe Ser Ser Asn Leu Asp Pro Ser Leu
195 200 205

Val Glu Gln Val Phe Leu Asp Lys Thr Leu Asn Ala Ser Phe His Trp
210 215 220

Leu Gly Ser Thr Tyr Gln Leu Val Asp Ile His Val Thr Glu Met Glu
225 230 235 240

Ser Ser Val Tyr Gln Pro Thr Ser Ser Ser Ser Thr Gln His Phe Tyr
245 250 255

Leu Asn Phe Thr Ile Thr Asn Leu Pro Tyr Ser Gln Asp Lys Ala Gln
260 265 270

Pro Gly Thr Thr Asn Tyr Gln Arg Asn Lys Arg Asn Ile Glu Asp Ala
275 280 285

Leu Asn Gln Leu Phe Arg Asn Ser Ser Ile Lys Ser Tyr Phe Ser Asp
 290 295 300
 Cys Gln Val Ser Thr Phe Arg Ser Val Pro Asn Arg His His Thr Gly
 305 310 315 320
 Val Asp Ser Leu Cys Asn Phe Ser Pro Leu Ala Arg Arg Val Asp Arg
 325 330 335
 Val Ala Ile Tyr Glu Glu Phe Leu Arg Met Thr Arg Asn Gly Thr Gln
 340 345 350
 Leu Gln Asn Phe Thr Leu Asp Arg Ser Ser Val Leu Val Asp Gly Tyr
 355 360 365
 Ser Pro Asn Arg Asn Glu Pro Leu Thr Gly Asn Ser Asp Leu Pro Phe
 370 375 380
 Trp Ala Val Ile Leu Ile Gly Leu Ala Gly Leu Leu Gly Leu Ile Thr
 385 390 395 400
 Cys Leu Ile Cys Gly Val Leu Val Thr Thr Arg Arg Arg Lys Lys Glu
 405 410 415
 Gly Glu Tyr Asn Val Gln Gln Cys Pro Gly Tyr Tyr Gln Ser His
 420 425 430
 Leu Asp Leu Glu Asp Leu Gln
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 <211> 1799
 <212> PRT
 <213> Homo sapiens

 <400> 149
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 Ser Pro Ala Ser Pro Lys Gly Leu His Thr Gly Gly Thr Lys Arg Met
 35 40 45
 Glu Thr Thr Thr Thr Ala Leu Lys Thr Thr Thr Thr Ala Leu Lys Thr
 50 55 60
 Thr Ser Arg Ala Thr Leu Thr Thr Ser Val Tyr Thr Pro Thr Leu Gly
 65 70 75 80
 Thr Leu Thr Pro Leu Asn Ala Ser Arg Gln Met Ala Ser Thr Ile Leu

85					90					95					
Thr	Glu	Met	Met	Ile	Thr	Thr	Pro	Tyr	Val	Phe	Pro	Asp	Val	Pro	Glu
			100					105					110		
Thr	Thr	Ser	Ser	Leu	Ala	Thr	Ser	Leu	Gly	Ala	Glu	Thr	Ser	Thr	Ala
			115					120					125		
Leu	Pro	Arg	Thr	Thr	Pro	Ser	Val	Leu	Asn	Arg	Glu	Ser	Glu	Thr	Thr
			130					135				140			
Ala	Ser	Leu	Val	Ser	Arg	Ser	Gly	Ala	Glu	Arg	Ser	Pro	Val	Ile	Gln
					150					155					160
Thr	Leu	Asp	Val	Ser	Ser	Ser	Glu	Pro	Asp	Thr	Thr	Ala	Ser	Trp	Val
				165					170					175	
Ile	His	Pro	Ala	Glu	Thr	Ile	Pro	Thr	Val	Ser	Lys	Thr	Thr	Pro	Asn
			180					185					190		
Phe	Phe	His	Ser	Glu	Leu	Asp	Thr	Val	Ser	Ser	Thr	Ala	Thr	Ser	His
			195				200					205			
Gly	Ala	Asp	Val	Ser	Ser	Ala	Ile	Pro	Thr	Asn	Ile	Ser	Pro	Ser	Glu
		210				215					220				
Leu	Asp	Ala	Leu	Thr	Pro	Leu	Val	Thr	Ile	Ser	Gly	Thr	Asp	Thr	Ser
	225				230						235				240
Thr	Thr	Phe	Pro	Thr	Leu	Thr	Lys	Ser	Pro	His	Glu	Thr	Glu	Thr	Arg
				245					250					255	
Thr	Thr	Trp	Leu	Thr	His	Pro	Ala	Glu	Thr	Ser	Ser	Thr	Ile	Pro	Arg
			260					265					270		
Thr	Ile	Pro	Asn	Phe	Ser	His	His	Glu	Ser	Asp	Ala	Thr	Pro	Ser	Ile
		275					280					285			
Ala	Thr	Ser	Pro	Gly	Ala	Glu	Thr	Ser	Ser	Ala	Ile	Pro	Ile	Met	Thr
			290			295					300				
Val	Ser	Pro	Gly	Ala	Glu	Asp	Leu	Val	Thr	Ser	Gln	Val	Thr	Ser	Ser
	305				310						315				320
Gly	Thr	Asp	Arg	Asn	Met	Thr	Ile	Pro	Thr	Leu	Thr	Leu	Ser	Pro	Gly
				325					330					335	
Glu	Pro	Lys	Thr	Ile	Ala	Ser	Leu	Val	Thr	His	Pro	Glu	Ala	Gln	Thr
			340					345					350		
Ser	Ser	Ala	Ile	Pro	Thr	Ser	Thr	Ile	Ser	Pro	Ala	Val	Ser	Arg	Leu
			355				360					365			
Val	Thr	Ser	Met	Val	Thr	Ser	Leu	Ala	Ala	Lys	Thr	Ser	Thr	Thr	Asn
			370			375					380				
Arg	Ala	Leu	Thr	Asn	Ser	Pro	Gly	Glu	Pro	Ala	Thr	Thr	Val	Ser	Leu
	385				390					395					400

Val Thr His Pro Ala Gln Thr Ser Pro Thr Val Pro Trp Thr Thr Ser
 405 410 415
 Ile Phe Phe His Ser Lys Ser Asp Thr Pro Ser Met Thr Thr Ser
 420 425 430
 His Gly Ala Glu Ser Ser Ser Ala Val Pro Thr Pro Thr Val Ser Thr
 435 440 445
 Glu Val Pro Gly Val Val Thr Pro Leu Val Thr Ser Ser Arg Ala Val
 450 455 460
 Ile Ser Thr Thr Ile Pro Ile Leu Thr Leu Ser Pro Gly Glu Pro Glu
 465 470 475 480
 Thr Thr Pro Ser Met Ala Thr Ser His Gly Glu Glu Ala Ser Ser Ala
 485 490 495
 Ile Pro Thr Pro Thr Val Ser Pro Gly Val Pro Gly Val Val Thr Ser
 500 505 510
 Leu Val Thr Ser Ser Arg Ala Val Thr Ser Thr Thr Ile Pro Ile Leu
 515 520 525
 Thr Phe Ser Leu Gly Glu Pro Glu Thr Thr Pro Ser Met Ala Thr Ser
 530 535 540
 His Gly Thr Glu Ala Gly Ser Ala Val Pro Thr Val Leu Pro Glu Val
 545 550 555 560
 Pro Gly Met Val Thr Ser Leu Val Ala Ser Ser Arg Ala Val Thr Ser
 565 570 575
 Thr Thr Leu Pro Thr Leu Thr Leu Ser Pro Gly Glu Pro Glu Thr Thr
 580 585 590
 Pro Ser Met Ala Thr Ser His Gly Ala Glu Ala Ser Ser Thr Val Pro
 595 600 605
 Thr Val Ser Pro Glu Val Pro Gly Val Val Thr Ser Leu Val Thr Ser
 610 615 620
 Ser Ser Gly Val Asn Ser Thr Ser Ile Pro Thr Leu Ile Leu Ser Pro
 625 630 635 640
 Gly Glu Leu Glu Thr Thr Pro Ser Met Ala Thr Ser His Gly Ala Glu
 645 650 655
 Ala Ser Ser Ala Val Pro Thr Pro Thr Val Ser Pro Gly Val Ser Gly
 660 665 670
 Val Val Thr Pro Leu Val Thr Ser Ser Arg Ala Val Thr Ser Thr Thr
 675 680 685
 Ile Pro Ile Leu Thr Leu Ser Ser Ser Glu Pro Glu Thr Thr Pro Ser
 690 695 700
 Met Ala Thr Ser His Gly Val Glu Ala Ser Ser Ala Val Leu Thr Val
 705 710 715 720

Ser Pro Glu Val Pro Gly Met Val Thr Ser Leu Val Thr Ser Ser Arg
 725 730 735
 Ala Val Thr Ser Thr Thr Ile Pro Thr Leu Thr Ile Ser Ser Asp Glu
 740 745 750
 Pro Glu Thr Thr Thr Ser Leu Val Thr His Ser Glu Ala Lys Met Ile
 755 760 765
 Ser Ala Ile Pro Thr Leu Ala Val Ser Pro Thr Val Gln Gly Leu Val
 770 775 780
 Thr Ser Leu Val Thr Ser Ser Gly Ser Glu Thr Ser Ala Phe Ser Asn
 785 790 795 800
 Leu Thr Val Ala Ser Ser Gln Pro Glu Thr Ile Asp Ser Trp Val Ala
 805 810 815
 His Pro Gly Thr Glu Ala Ser Ser Val Val Pro Thr Leu Thr Val Ser
 820 825 830
 Thr Gly Glu Pro Phe Thr Asn Ile Ser Leu Val Thr His Pro Ala Glu
 835 840 845
 Ser Ser Ser Thr Leu Pro Arg Thr Thr Ser Arg Phe Ser His Ser Glu
 850 855 860
 Leu Asp Thr Met Pro Ser Thr Val Thr Ser Pro Glu Ala Glu Ser Ser
 865 870 875 880
 Ser Ala Ile Ser Thr Thr Ile Ser Pro Gly Ile Pro Gly Val Leu Thr
 885 890 895
 Ser Leu Val Thr Ser Ser Gly Arg Asp Ile Ser Ala Thr Phe Pro Thr
 900 905 910
 Val Pro Glu Ser Pro His Glu Ser Glu Ala Thr Ala Ser Trp Val Thr
 915 920 925
 His Pro Ala Val Thr Ser Thr Thr Val Pro Arg Thr Thr Pro Asn Tyr
 930 935 940
 Ser His Ser Glu Pro Asp Thr Thr Pro Ser Ile Ala Thr Ser Pro Gly
 945 950 955 960
 Ala Glu Ala Thr Ser Asp Phe Pro Thr Ile Thr Val Ser Pro Asp Val
 965 970 975
 Pro Asp Met Val Thr Ser Gln Val Thr Ser Ser Gly Thr Asp Thr Ser
 980 985 990
 Ile Thr Ile Pro Thr Leu Thr Leu Ser Ser Gly Glu Pro Glu Thr Thr
 995 1000 1005
 Thr Ser Phe Ile Thr Tyr Ser Glu Thr His Thr Ser Ser Ala Ile
 1010 1015 1020
 Pro Thr Leu Pro Val Ser Pro Gly Ala Ser Lys Met Leu Thr Ser

1025	1030	1035
Leu Val Ile Ser Ser Gly Thr	Asp Ser Thr Thr Thr	Phe Pro Thr
1040	1045	1050
Leu Thr Glu Thr Pro Tyr Glu	Pro Glu Thr Thr Ala	Ile Gln Leu
1055	1060	1065
Ile His Pro Ala Glu Thr Asn	Thr Met Val Pro Arg	Thr Thr Pro
1070	1075	1080
Lys Phe Ser His Ser Lys Ser	Asp Thr Thr Leu Pro	Val Ala Ile
1085	1090	1095
Thr Ser Pro Gly Pro Glu Ala	Ser Ser Ala Val Ser	Thr Thr Thr
1100	1105	1110
Ile Ser Pro Asp Met Ser Asp	Leu Val Thr Ser Leu	Val Pro Ser
1115	1120	1125
Ser Gly Thr Asp Thr Ser Thr	Thr Phe Pro Thr Leu	Ser Glu Thr
1130	1135	1140
Pro Tyr Glu Pro Glu Thr Thr	Ala Thr Trp Leu Thr	His Pro Ala
1145	1150	1155
Glu Thr Ser Thr Thr Val Ser	Gly Thr Ile Pro Asn	Phe Ser His
1160	1165	1170
Arg Gly Ser Asp Thr Ala Pro	Ser Met Val Thr Ser	Pro Gly Val
1175	1180	1185
Asp Thr Arg Ser Gly Val Pro	Thr Thr Thr Ile Pro	Pro Ser Ile
1190	1195	1200
Pro Gly Val Val Thr Ser Gln	Val Thr Ser Ser Ala	Thr Asp Thr
1205	1210	1215
Ser Thr Ala Ile Pro Thr Leu	Thr Pro Ser Pro Gly	Glu Pro Glu
1220	1225	1230
Thr Thr Ala Ser Ser Ala Thr	His Pro Gly Thr Gln	Thr Gly Phe
1235	1240	1245
Thr Val Pro Ile Arg Thr Val	Pro Ser Ser Glu Pro	Asp Thr Met
1250	1255	1260
Ala Ser Trp Val Thr His Pro	Pro Gln Thr Ser Thr	Pro Val Ser
1265	1270	1275
Arg Thr Thr Ser Ser Phe Ser	His Ser Ser Pro Asp	Ala Thr Pro
1280	1285	1290
Val Met Ala Thr Ser Pro Arg	Thr Glu Ala Ser Ser	Ala Val Leu
1295	1300	1305
Thr Thr Ile Ser Pro Gly Ala	Pro Glu Met Val Thr	Ser Gln Ile
1310	1315	1320

Thr Ser Ser Gly Ala Ala Thr Ser Thr Thr Val Pro Thr Leu Thr
 1325 1330 1335
 His Ser Pro Gly Met Pro Glu Thr Thr Ala Leu Leu Ser Thr His
 1340 1345 1350
 Pro Arg Thr Glu Thr Ser Lys Thr Phe Pro Ala Ser Thr Val Phe
 1355 1360 1365
 Pro Gln Val Ser Glu Thr Thr Ala Ser Leu Thr Ile Arg Pro Gly
 1370 1375 1380
 Ala Glu Thr Ser Thr Ala Leu Pro Thr Gln Thr Thr Ser Ser Leu
 1385 1390 1395
 Phe Thr Leu Leu Val Thr Gly Thr Ser Arg Val Asp Leu Ser Pro
 1400 1405 1410
 Thr Ala Ser Pro Gly Val Ser Ala Lys Thr Ala Pro Leu Ser Thr
 1415 1420 1425
 His Pro Gly Thr Glu Thr Ser Thr Met Ile Pro Thr Ser Thr Leu
 1430 1435 1440
 Ser Leu Gly Leu Leu Glu Thr Thr Gly Leu Leu Ala Thr Ser Ser
 1445 1450 1455
 Ser Ala Glu Thr Ser Thr Ser Thr Leu Thr Leu Thr Val Ser Pro
 1460 1465 1470
 Ala Val Ser Gly Leu Ser Ser Ala Ser Ile Thr Thr Asp Lys Pro
 1475 1480 1485
 Gln Thr Val Thr Ser Trp Asn Thr Glu Thr Ser Pro Ser Val Thr
 1490 1495 1500
 Ser Val Gly Pro Pro Glu Phe Ser Arg Thr Val Thr Gly Thr Thr
 1505 1510 1515
 Met Thr Leu Ile Pro Ser Glu Met Pro Thr Pro Pro Lys Thr Ser
 1520 1525 1530
 His Gly Glu Gly Val Ser Pro Thr Thr Ile Leu Arg Thr Thr Met
 1535 1540 1545
 Val Glu Ala Thr Asn Leu Ala Thr Thr Gly Ser Ser Pro Thr Val
 1550 1555 1560
 Ala Lys Thr Thr Thr Thr Phe Asn Thr Leu Ala Gly Ser Leu Phe
 1565 1570 1575
 Thr Pro Leu Thr Thr Pro Gly Met Ser Thr Leu Ala Ser Glu Ser
 1580 1585 1590
 Val Thr Ser Arg Thr Ser Tyr Asn His Arg Ser Trp Ile Ser Thr
 1595 1600 1605
 Thr Ser Ser Tyr Asn Arg Arg Tyr Trp Thr Pro Ala Thr Ser Thr
 1610 1615 1620

Pro Val Thr Ser Thr Phe Ser Pro Gly Ile Ser Thr Ser Ser Ile
1625 1630 1635

Pro Ser Ser Thr Ala Ala Thr Val Pro Phe Met Val Pro Phe Thr
1640 1645 1650

Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg
1655 1660 1665

His Pro Gly Ser Arg Lys Phe Asn Ala Thr Glu Arg Glu Leu Gln
1670 1675 1680

Gly Leu Leu Lys Pro Leu Phe Arg Asn Ser Ser Leu Glu Tyr Leu
1685 1690 1695

Tyr Ser Gly Cys Arg Leu Ala Ser Leu Arg Pro Glu Lys Asp Ser
1700 1705 1710

Ser Ala Met Ala Val Asp Ala Ile Cys Thr His Arg Pro Asp Pro
1715 1720 1725

Glu Asp Leu Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser
1730 1735 1740

Asn Leu Thr Asn Gly Ile Gln Glu Leu Gly Pro Tyr Thr Leu Asp
1745 1750 1755

Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Met
1760 1765 1770

Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Val Gly Thr
1775 1780 1785

Ser Gly Thr Pro Ser Ser Ser Pro Ser Pro Thr
1790 1795

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<211> 156

<212> PRT

<213> Homo sapiens

<400> 150

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Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys
35 40 45

Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu

50 55 60
 Arg Pro Glu Lys His Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr
 65 70 75 80
 His Arg Val Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr
 85 90 95
 Trp Glu Leu Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr
 100 105 110
 Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Asn Pro Arg Ser
 115 120 125
 Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala
 130 135 140
 Thr Ser Gly Thr Pro Ser Ser Leu Pro Lys Leu Thr
 145 150 155

<210> 151

<211> 507

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (1)..(507)

<400> 151
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 Glu Pro Gly Pro Leu Leu Ile Pro Phe Thr Phe Asn Phe Thr Ile Thr
 20 25 30
 aac ctg cat tat gag gaa aac atg caa cac cct ggt tcc agg aag ttc 144
 Asn Leu His Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe
 35 40 45
 aac acc acg gag agg gtt ctg cag ggt ctg ctc aag ccc ttg ttc aag 192
 Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys
 50 55 60
 aac acc agt gtt ggc cct ctg tac tct ggc tgc aga ctg acc ttg ctc 240
 Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu
 65 70 75 80

aga cct gag aag cat gag gca gcc act gga gtg gac acc atc tgt acc 288
 Arg Pro Glu Lys His Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr
 85 90 95

cac cgc gtt gat ccc atc gga cct gga ctg gac aga gag cgg cta tac 336
 His Arg Val Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr
 100 105 110

tgg gag ctg agc cag ctg acc aac agc atc aca gag ctg gga ccc tac 384
 Trp Glu Leu Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr
 115 120 125

acc ctg gac agg gac agt ctc tat gtc aat ggc ttc aac cct cgg agc 432
 Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Asn Pro Arg Ser
 130 135 140

tct gtg cca acc acc agc act cct ggg acc tcc aca gtg cac ctg gca 480
 Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala
 145 150 155 160

acc tct ggg act cca tcc tcc ctg cct 507
 Thr Ser Gly Thr Pro Ser Ser Leu Pro
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<210> 152
 <211> 169
 <212> PRT
 <213> Homo sapiens

<400> 152

Met Arg Gly Ser His His His His His His Gly Ser Met Gly His Thr
 1 5 10 15

Glu Pro Gly Pro Leu Leu Ile Pro Phe Thr Phe Asn Phe Thr Ile Thr
 20 25 30

Asn Leu His Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe
 35 40 45

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys
 50 55 60

Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu
 65 70 75 80

Arg Pro Glu Lys His Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr
 85 90 95

<212> PRT

<213> Homo sapiens

<400> 154

Arg Leu Tyr Trp Glu Leu Ser Gln Leu
1 5

<210> 155

<211> 9

<212> PRT

<213> Homo sapiens

<400> 155

Thr Leu Asp Arg Asp Ser Leu Tyr Val
1 5

<210> 156

<211> 9

<212> PRT

<213> Homo sapiens

<400> 156

Val Leu Gln Gly Leu Leu Lys Pro Leu
1 5

<210> 157

<211> 9

<212> PRT

<213> Homo sapiens

<400> 157

Gln Leu Thr Asn Ser Ile Thr Glu Leu
1 5

<210> 158

<211> 780

<212> PRT

<213> Homo sapiens

<400> 158

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Ala Thr Val Pro Phe Met Val Pro Phe Thr Leu Asn Phe Thr Ile Thr
1          5          10          15

Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe
          20          25          30

Asn Ala Thr Glu Arg Glu Leu Gln Gly Leu Leu Lys Pro Leu Phe Arg
          35          40          45

Asn Ser Ser Leu Glu Tyr Leu Tyr Ser Gly Cys Arg Leu Ala Ser Leu
          50          55          60

Arg Pro Glu Lys Asp Ser Ser Ala Met Ala Val Asp Ala Ile Cys Thr
          65          70          75          80

His Arg Pro Asp Pro Glu Asp Leu Gly Leu Asp Arg Glu Arg Leu Tyr
          85          90          95

Trp Glu Leu Ser Asn Leu Thr Asn Gly Ile Gln Glu Leu Gly Pro Tyr
          100          105          110

Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser
          115          120          125

Ser Met Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Val Gly
          130          135          140

Thr Ser Gly Thr Pro Ser Ser Ser Pro Ser Pro Thr Ala Ala Gly Pro
          145          150          155          160

Leu Leu Met Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr
          165          170          175

Glu Glu Asp Met Arg Arg Thr Gly Ser Arg Lys Phe Asn Thr Met Glu
          180          185          190

Ser Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val
          195          200          205

Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys
          210          215          220

Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu Asp
          225          230          235          240

Pro Lys Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp Glu Leu Ser
          245          250          255

Lys Leu Thr Asn Asp Ile Glu Glu Leu Gly Pro Tyr Thr Leu Asp Arg
          260          265          270

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Asn Ser Leu Tyr Val Asn Gly Phe Thr His Gln Ser Ser Val Ser Thr
 275 280
 Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Arg Thr Ser Gly Thr
 290 295 300
 Pro Ser Ser Leu Ser Ser Pro Thr Ile Met Ala Gly Pro Leu Leu Val
 305 310 315 320
 Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Gly Glu Asp
 325 330 335
 Met Gly His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu
 340 345 350
 Gln Gly Leu Leu Gly Pro Ile Phe Lys Asn Thr Ser Val Gly Pro Leu
 355 360 365
 Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Ser Glu Lys Asp Gly Ala
 370 375 380
 Ala Thr Gly Val Asp Ala Ile Cys Ile His His Leu Asp Pro Lys Ser
 385 390 395 400
 Pro Gly Leu Asn Arg Glu Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr
 405 410 415
 Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu
 420 425 430
 Tyr Val Asn Gly Phe Thr His Arg Thr Ser Val Pro Thr Ser Ser Thr
 435 440 445
 Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Phe Ser
 450 455 460
 Leu Pro Ser Pro Ala Thr Ala Gly Pro Leu Leu Val Leu Phe Thr Leu
 465 470 475 480
 Asn Phe Thr Ile Thr Asn Leu Lys Tyr Glu Glu Asp Met His Arg Pro
 485 490 495
 Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Thr Leu Leu
 500 505 510
 Gly Pro Met Phe Lys Asn Thr Ser Val Gly Leu Leu Tyr Ser Gly Cys
 515 520 525
 Arg Leu Thr Leu Leu Arg Ser Glu Lys Asp Gly Ala Ala Thr Gly Val
 530 535 540
 Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser Pro Gly Leu Asp
 545 550 555 560
 Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Gly Ile Lys
 565 570 575
 Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly
 580 585 590

Phe Thr His Trp Ile Pro Val Pro Thr Ser Ser Thr Pro Gly Thr Ser
 595 600 605
 Thr Val Asp Leu Gly Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr
 610 615 620
 Ala Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr
 625 630 635 640
 Asn Leu Gln Tyr Glu Glu Asp Met His His Pro Gly Ser Arg Lys Phe
 645 650 655
 Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Met Phe Lys
 660 665 670
 Asn Thr Ser Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu
 675 680 685
 Arg Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr
 690 695 700
 His Arg Leu Asp Pro Lys Ser Pro Gly Val Asp Arg Glu Gln Leu Tyr
 705 710 715 720
 Trp Glu Leu Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr
 725 730 735
 Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Gln Thr
 740 745 750
 Ser Ala Pro Asn Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly
 755 760 765
 Thr Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr
 770 775 780

<210> 159

<211> 780

<212> PRT

<213> Homo sapiens

<400> 159

Ser Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr
 1 5 10 15
 Asn Leu Gln Tyr Glu Glu Asp Met His His Pro Gly Ser Arg Lys Phe
 20 25 30
 Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Met Phe Lys
 35 40 45
 Asn Thr Ser Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu

50	55	60
Arg Pro Glu Lys Asn Gly Ala Ala Thr Gly Met Asp Ala Ile Cys Ser 65 70 75 80		
His Arg Leu Asp Pro Lys Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr 85 90 95		
Trp Glu Leu Ser Gln Leu Thr His Gly Ile Lys Glu Leu Gly Pro Tyr 100 105 110		
Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser 115 120 125		
Ser Val Ala Pro Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly 130 135 140		
Thr Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr Thr Ala Val Pro 145 150 155 160		
Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr 165 170 175		
Gly Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu 180 185 190		
Arg Val Leu Gln Gly Leu Leu Gly Pro Leu Phe Lys Asn Ser Ser Val 195 200 205		
Gly Pro Leu Tyr Ser Gly Cys Arg Leu Ile Ser Leu Arg Ser Glu Lys 210 215 220		
Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His His Leu Asn 225 230 235 240		
Pro Gln Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Gln Leu Ser 245 250 255		
Gln Met Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg 260 265 270		
Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Gly Leu Thr 275 280 285		
Thr Ser Thr Pro Trp Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr 290 295 300		
Pro Ser Pro Val Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Val Pro 305 310 315 320		
Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met 325 330 335		
His Arg Pro Gly Ser Arg Lys Phe Asn Ala Thr Glu Arg Val Leu Gln 340 345 350		
Gly Leu Leu Ser Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu Tyr 355 360 365		

Ser	Gly	Cys	Arg	Leu	Thr	Ser	Leu	Arg	Pro	Glu	Lys	Asp	Gly	Ala	Ala
370						375					380				
Thr	Gly	Met	Asp	Ala	Val	Cys	Leu	Tyr	His	Pro	Asn	Pro	Lys	Arg	Pro
385					390					395					400
Gly	Leu	Asp	Arg	Glu	Gln	Leu	Tyr	Trp	Glu	Leu	Ser	Gln	Leu	Thr	His
				405					410					415	
Asn	Ile	Thr	Glu	Leu	Gly	Pro	Tyr	Ser	Leu	Asp	Arg	Asp	Ser	Leu	Tyr
			420					425					430		
Val	Asn	Gly	Phe	Thr	His	Gln	Asn	Ser	Val	Pro	Thr	Thr	Ser	Thr	Pro
			435				440					445			
Gly	Thr	Ser	Thr	Val	Tyr	Trp	Ala	Thr	Thr	Gly	Thr	Pro	Ser	Ser	Phe
						455					460				
Pro	Gly	His	Thr	Glu	Pro	Gly	Pro	Leu	Leu	Ile	Pro	Phe	Thr	Phe	Asn
465					470					475					480
Phe	Thr	Ile	Thr	Asn	Leu	His	Tyr	Glu	Glu	Asn	Met	Gln	His	Pro	Gly
				485					490					495	
Ser	Arg	Lys	Phe	Asn	Thr	Thr	Glu	Arg	Val	Leu	Gln	Gly	Leu	Leu	Lys
			500					505					510		
Pro	Leu	Phe	Lys	Asn	Thr	Ser	Val	Gly	Pro	Leu	Tyr	Ser	Gly	Cys	Arg
			515				520					525			
Leu	Thr	Ser	Leu	Arg	Pro	Glu	Lys	Asp	Gly	Ala	Ala	Thr	Gly	Met	Asp
						535					540				
Ala	Val	Cys	Leu	Tyr	His	Pro	Asn	Pro	Lys	Arg	Pro	Gly	Leu	Asp	Arg
545					550					555					560
Glu	Gln	Leu	Tyr	Cys	Glu	Leu	Ser	Gln	Leu	Thr	His	Asn	Ile	Thr	Glu
				565					570					575	
Leu	Gly	Pro	Tyr	Ser	Leu	Asp	Arg	Asp	Ser	Leu	Tyr	Val	Asn	Gly	Phe
				580				585					590		
Thr	His	Gln	Asn	Ser	Val	Pro	Thr	Thr	Ser	Thr	Pro	Gly	Thr	Ser	Thr
			595				600						605		
Val	Tyr	Trp	Ala	Thr	Thr	Gly	Thr	Pro	Ser	Ser	Phe	Pro	Gly	His	Thr
						615					620				
Glu	Pro	Gly	Pro	Leu	Leu	Ile	Pro	Phe	Thr	Phe	Asn	Phe	Thr	Ile	Thr
625					630					635					640
Asn	Leu	His	Tyr	Glu	Glu	Asn	Met	Gln	His	Pro	Gly	Ser	Arg	Lys	Phe
				645					650					655	
Asn	Thr	Thr	Glu	Arg	Val	Leu	Gln	Gly	Leu	Leu	Lys	Pro	Leu	Phe	Lys
				660				665					670		
Asn	Thr	Ser	Val	Gly	Pro	Leu	Tyr	Ser	Gly	Cys	Arg	Leu	Thr	Leu	Leu
							680					685			

Arg Pro Glu Lys His Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr
690 695 700

His Arg Val Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr
705 710 715 720

Trp Glu Leu Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr
725 730 735

Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Asn Pro Arg Ser
740 745 750

Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala
755 760 765

Thr Ser Gly Thr Pro Ser Ser Leu Pro Gly His Thr
770 775 780

<210> 160

<211> 624

<212> PRT

<213> Homo sapiens

<400> 160

Thr Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr
1 5 10 15

Asn Leu Gln Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Arg Phe
20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Thr Pro Leu Phe Lys
35 40 45

Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu
50 55 60

Arg Pro Glu Lys Gln Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr
65 70 75 80

His Arg Val Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr
85 90 95

Trp Glu Leu Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr
100 105 110

Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Asn Pro Trp Ser
115 120 125

Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala
130 135 140

Thr Ser Gly Thr Pro Ser Ser Leu Pro Gly His Thr Ala Pro Val Pro

145 150 155 160
 Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr Asp Leu His Tyr
 165 170 175
 Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu
 180 185 190
 Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Val
 195 200 205
 Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys
 210 215 220
 His Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr Leu Arg Leu Asp
 225 230 235 240
 Pro Thr Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser
 245 250 255
 Gln Leu Thr Asn Ser Val Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg
 260 265 270
 Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Pro Thr
 275 280 285
 Thr Ser Ile Pro Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly Thr
 290 295 300
 Pro Ala Ser Leu Pro Gly His Thr Ala Pro Gly Pro Leu Leu Val Pro
 305 310 315 320
 Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met
 325 330 335
 Arg His Pro Gly Ser Arg Lys Phe Ser Thr Thr Glu Arg Val Leu Gln
 340 345 350
 Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu Tyr
 355 360 365
 Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala Ala
 370 375 380
 Thr Arg Val Asp Ala Val Cys Thr His Arg Pro Asp Pro Lys Ser Pro
 385 390 395 400
 Gly Leu Asp Arg Glu Arg Leu Tyr Trp Lys Leu Ser Gln Leu Thr His
 405 410 415
 Gly Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg His Ser Leu Tyr
 420 425 430
 Val Asn Gly Phe Thr His Gln Ser Ser Met Thr Thr Thr Arg Thr Pro
 435 440 445
 Asp Thr Ser Thr Met His Leu Ala Thr Ser Arg Thr Pro Ala Ser Leu
 450 455 460

Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu Val Leu Phe Thr Ile Asn
 465 470 475 480
 Phe Thr Ile Thr Asn Gln Arg Tyr Glu Glu Asn Met His His Pro Gly
 485 490 495
 Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Arg
 500 505 510
 Pro Val Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg
 515 520 525
 Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala Ala Thr Lys Val Asp
 530 535 540
 Ala Ile Cys Thr Tyr Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg
 545 550 555 560
 Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His Ser Ile Thr Glu
 565 570 575
 Leu Gly Pro Tyr Thr Gln Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe
 580 585 590
 Thr His Arg Ser Ser Val Pro Thr Thr Ser Ile Pro Gly Thr Ser Ala
 595 600 605
 Val His Leu Glu Thr Ser Gly Thr Pro Ala Ser Leu Pro Gly His Thr
 610 615 620
 <210> 161
 <211> 468
 <212> PRT
 <213> Homo sapiens

 <400> 161
 Ala Thr Gly Pro Val Leu Leu Pro Phe Thr Leu Asn Phe Thr Ile Thr
 1 5 10 15
 Asn Leu Gln Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe
 20 25 30
 Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Met Pro Leu Phe Lys
 35 40 45
 Asn Thr Ser Val Ser Ser Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu
 50 55 60
 Arg Pro Glu Lys Asp Gly Ala Ala Thr Arg Val Asp Ala Val Cys Thr
 65 70 75 80
 His Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Arg Leu Tyr
 85 90 95

Trp Lys Leu Ser Gln Leu Thr His Gly Ile Thr Glu Leu Gly Pro Tyr
 100 105 110
 Thr Leu Asp Arg His Ser Leu Tyr Val Asn Gly Phe Thr His Gln Ser
 115 120 125
 Ser Met Thr Thr Thr Arg Thr Pro Asp Thr Ser Thr Met His Leu Ala
 130 135 140
 Thr Ser Arg Thr Pro Ala Ser Leu Ser Gly Pro Thr Thr Ala Ser Pro
 145 150 155 160
 Leu Leu Val Leu Phe Thr Ile Asn Phe Thr Ile Thr Asn Leu Arg Tyr
 165 170 175
 Glu Glu Asn Met His His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu
 180 185 190
 Arg Val Leu Gln Gly Leu Leu Arg Pro Val Phe Lys Asn Thr Ser Val
 195 200 205
 Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys Lys
 210 215 220
 Asp Gly Ala Ala Thr Lys Val Asp Ala Ile Cys Thr Tyr Arg Pro Asp
 225 230 235 240
 Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser
 245 250 255
 Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Gln Asp Arg
 260 265 270
 Asp Ser Leu Tyr Asn Val Gly Phe Thr Gln Arg Ser Ser Val Pro Thr
 275 280 285
 Thr Ser Val Pro Gly Thr Pro Thr Val Asp Leu Gly Thr Ser Gly Thr
 290 295 300
 Pro Val Ser Lys Pro Gly Pro Ser Ala Ala Ser Pro Leu Leu Val Leu
 305 310 315 320
 Phe Thr Leu Asn Gly Thr Ile Thr Asn Leu Arg Tyr Glu Glu Asn Met
 325 330 335
 Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln
 340 345 350
 Gly Leu Leu Arg Ser Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr
 355 360 365
 Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Thr Ala
 370 375 380
 Thr Gly Val Asp Ala Ile Cys Thr His His Pro Asp Pro Lys Ser Pro
 385 390 395 400
 Arg Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His

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                                405                                410                                415
Asn Ile Thr Glu Leu Gly His Tyr Ala Leu Asp Asn Asp Ser Leu Phe
                                420                                425                                430
Val Asn Gly Phe Thr His Arg Ser Ser Val Ser Thr Thr Ser Thr Pro
                                435                                440                                445
Gly Thr Pro Thr Val Tyr Leu Gly Ala Ser Lys Thr Pro Ala Ser Ile
                                450                                455                                460
Phe Gly Pro Ser
465

<210> 162

<211> 11721

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (1)..(11721)

<223> any x = any amino acid

<400> 162

Met Glu His Ile Thr Lys Ile Pro Asn Glu Ala Ala His Arg Gly Thr
1 5 10 15
Ile Arg Pro Val Lys Gly Pro Gln Thr Ser Thr Ser Pro Ala Ser Pro
20 25 30
Lys Gly Leu His Thr Gly Gly Thr Lys Arg Met Glu Thr Thr Thr Thr
35 40 45
Ala Leu Lys Thr Thr Thr Thr Ala Leu Lys Thr Thr Ser Arg Ala Thr
50 55 60
Leu Thr Thr Ser Val Tyr Thr Pro Thr Leu Gly Thr Leu Thr Pro Leu
65 70 75 80
Asn Ala Ser Arg Gln Met Ala Ser Thr Ile Leu Thr Glu Met Met Ile
85 90 95
Thr Thr Pro Tyr Val Phe Pro Asp Val Pro Glu Thr Thr Ser Ser Leu
100 105 110
Ala Thr Ser Leu Gly Ala Glu Thr Ser Thr Ala Leu Pro Arg Thr Thr
115 120 125

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Pro Ser Val Leu Asn Arg Glu Ser Glu Thr Thr Ala Ser Leu Val Ser
 130 135 140
 Arg Ser Gly Ala Glu Arg Ser Pro Val Ile Gln Thr Leu Asp Val Ser
 145 150 155 160
 Ser Ser Glu Pro Asp Thr Thr Ala Ser Trp Val Ile His Pro Ala Glu
 165 170 175
 Thr Ile Pro Thr Val Ser Lys Thr Thr Pro Asn Phe Phe His Ser Glu
 180 185 190
 Leu Asp Thr Val Ser Ser Thr Ala Thr Ser His Gly Ala Asp Val Ser
 195 200 205
 Ser Ala Ile Pro Thr Asn Ile Ser Pro Ser Glu Leu Asp Ala Leu Thr
 210 215 220
 Pro Leu Val Thr Ile Ser Gly Thr Asp Thr Ser Thr Thr Phe Pro Thr
 225 230 235 240
 Leu Thr Lys Ser Pro His Glu Thr Glu Thr Arg Thr Thr Trp Leu Thr
 245 250 255
 His Pro Ala Glu Thr Ser Ser Thr Ile Pro Arg Thr Ile Pro Asn Phe
 260 265 270
 Ser His His Glu Ser Asp Ala Thr Pro Ser Ile Ala Thr Ser Pro Gly
 275 280 285
 Ala Glu Thr Ser Ser Ala Ile Pro Ile Met Thr Val Ser Pro Gly Ala
 290 295 300
 Glu Asp Leu Val Thr Ser Gln Val Thr Ser Ser Gly Thr Asp Arg Asn
 305 310 315 320
 Met Thr Ile Pro Thr Leu Thr Leu Ser Pro Gly Glu Pro Lys Thr Ile
 325 330 335
 Ala Ser Leu Val Thr His Pro Glu Ala Gln Thr Ser Ser Ala Ile Pro
 340 345 350
 Thr Ser Thr Ile Ser Pro Ala Val Ser Arg Leu Val Thr Ser Met Val
 355 360 365
 Thr Ser Leu Ala Ala Lys Thr Ser Thr Thr Asn Arg Ala Leu Thr Asn
 370 375 380
 Ser Pro Gly Glu Pro Ala Thr Thr Val Ser Leu Val Thr His Pro Ala
 385 390 395 400
 Gln Thr Ser Pro Thr Val Pro Trp Thr Thr Ser Ile Phe Phe His Ser
 405 410 415
 Lys Ser Asp Thr Thr Pro Ser Met Thr Thr Ser His Gly Ala Glu Ser
 420 425 430
 Ser Ser Ala Val Pro Thr Pro Thr Val Ser Thr Glu Val Pro Gly Val

435		440		445
Val Thr Pro Leu Val Thr Ser Ser Arg Ala Val Ile Ser Thr Thr Ile				
450		455		460
Pro Ile Leu Thr Leu Ser Pro Gly Glu Pro Glu Thr Thr Pro Ser Met				
465		470		480
Ala Thr Ser His Gly Glu Glu Ala Ser Ser Ala Ile Pro Thr Pro Thr				
	485		490	495
Val Ser Pro Gly Val Pro Gly Val Val Thr Ser Leu Val Thr Ser Ser				
	500		505	510
Arg Ala Val Thr Ser Thr Thr Ile Pro Ile Leu Thr Phe Ser Leu Gly				
	515		520	525
Glu Pro Glu Thr Thr Pro Ser Met Ala Thr Ser His Gly Thr Glu Ala				
	530		535	540
Gly Ser Ala Val Pro Thr Val Leu Pro Glu Val Pro Gly Met Val Thr				
545		550		555
Ser Leu Val Ala Ser Ser Arg Ala Val Thr Ser Thr Thr Leu Pro Thr				
	565		570	575
Leu Thr Leu Ser Pro Gly Glu Pro Glu Thr Thr Pro Ser Met Ala Thr				
	580		585	590
Ser His Gly Ala Glu Ala Ser Ser Thr Val Pro Thr Val Ser Pro Glu				
	595		600	605
Val Pro Gly Val Val Thr Ser Leu Val Thr Ser Ser Ser Gly Val Asn				
	610		615	620
Ser Thr Ser Ile Pro Thr Leu Ile Leu Ser Pro Gly Glu Leu Glu Thr				
625		630		635
Thr Pro Ser Met Ala Thr Ser His Gly Ala Glu Ala Ser Ser Ala Val				
	645		650	655
Pro Thr Pro Thr Val Ser Pro Gly Val Ser Gly Val Val Thr Pro Leu				
	660		665	670
Val Thr Ser Ser Arg Ala Val Thr Ser Thr Thr Ile Pro Ile Leu Thr				
	675		680	685
Leu Ser Ser Ser Glu Pro Glu Thr Thr Pro Ser Met Ala Thr Ser His				
	690		695	700
Gly Val Glu Ala Ser Ser Ala Val Leu Thr Val Ser Pro Glu Val Pro				
705		710		715
Gly Met Val Thr Ser Leu Val Thr Ser Ser Arg Ala Val Thr Ser Thr				
	725		730	735
Thr Ile Pro Thr Leu Thr Ile Ser Ser Asp Glu Pro Glu Thr Thr Thr				
	740		745	750

Ser Leu Val Thr His Ser Glu Ala Lys Met Ile Ser Ala Ile Pro Thr
 755 760 765
 Leu Ala Val Ser Pro Thr Val Gln Gly Leu Val Thr Ser Leu Val Thr
 770 775 780
 Ser Ser Gly Ser Glu Thr Ser Ala Phe Ser Asn Leu Thr Val Ala Ser
 785 790 795 800
 Ser Gln Pro Glu Thr Ile Asp Ser Trp Val Ala His Pro Gly Thr Glu
 805 810 815
 Ala Ser Ser Val Val Pro Thr Leu Thr Val Ser Thr Gly Glu Pro Phe
 820 825 830
 Thr Asn Ile Ser Leu Val Thr His Pro Ala Glu Ser Ser Ser Thr Leu
 835 840 845
 Pro Arg Thr Thr Ser Arg Phe Ser His Ser Glu Leu Asp Thr Met Pro
 850 855 860
 Ser Thr Val Thr Ser Pro Glu Ala Glu Ser Ser Ala Ile Ser Thr
 865 870 875 880
 Thr Ile Ser Pro Gly Ile Pro Gly Val Leu Thr Ser Leu Val Thr Ser
 885 890 895
 Ser Gly Arg Asp Ile Ser Ala Thr Phe Pro Thr Val Pro Glu Ser Pro
 900 905 910
 His Glu Ser Glu Ala Thr Ala Ser Trp Val Thr His Pro Ala Val Thr
 915 920 925
 Ser Thr Thr Val Pro Arg Thr Thr Pro Asn Tyr Ser His Ser Glu Pro
 930 935 940
 Asp Thr Thr Pro Ser Ile Ala Thr Ser Pro Gly Ala Glu Ala Thr Ser
 945 950 955 960
 Asp Phe Pro Thr Ile Thr Val Ser Pro Asp Val Pro Asp Met Val Thr
 965 970 975
 Ser Gln Val Thr Ser Ser Gly Thr Asp Thr Ser Ile Thr Ile Pro Thr
 980 985 990
 Leu Thr Leu Ser Ser Gly Glu Pro Glu Thr Thr Thr Ser Phe Ile Thr
 995 1000 1005
 Tyr Ser Glu Thr His Thr Ser Ser Ala Ile Pro Thr Leu Pro Val
 1010 1015 1020
 Ser Pro Gly Ala Ser Lys Met Leu Thr Ser Leu Val Ile Ser Ser
 1025 1030 1035
 Gly Thr Asp Ser Thr Thr Phe Pro Thr Leu Thr Glu Thr Pro
 1040 1045 1050
 Tyr Glu Pro Glu Thr Thr Ala Ile Gln Leu Ile His Pro Ala Glu
 1055 1060 1065

Thr Asn	Thr Met	Val Pro	Arg	Thr Thr	Pro Lys	Phe	Ser His	Ser	
1070			1075			1080			
Lys Ser	Asp Thr	Thr Leu	Pro	Val Ala	Ile Thr	Ser	Pro Gly	Pro	
1085			1090			1095			
Glu Ala	Ser Ser	Ala Val	Ser	Thr Thr	Thr Ile	Ser	Pro Asp	Met	
1100			1105			1110			
Ser Asp	Leu Val	Thr Ser	Leu	Val Pro	Ser Ser	Gly	Thr Asp	Thr	
1115			1120			1125			
Ser Thr	Thr Phe	Pro Thr	Leu	Ser Glu	Thr Pro	Tyr	Glu Pro	Glu	
1130			1135			1140			
Thr Thr	Ala Thr	Trp Leu	Thr	His Pro	Ala Glu	Thr	Ser Thr	Thr	
1145			1150			1155			
Val Ser	Gly Thr	Ile Pro	Asn	Phe Ser	His Arg	Gly	Ser Asp	Thr	
1160			1165			1170			
Ala Pro	Ser Met	Val Thr	Ser	Pro Gly	Val Asp	Thr	Arg Ser	Gly	
1175			1180			1185			
Val Pro	Thr Thr	Thr Ile	Pro	Pro Ser	Ile Pro	Gly	Val Val	Thr	
1190			1195			1200			
Ser Gln	Val Thr	Ser Ser	Ala	Thr Asp	Thr Ser	Thr	Ala Ile	Pro	
1205			1210			1215			
Thr Leu	Thr Pro	Ser Pro	Gly	Glu Pro	Glu Thr	Thr	Ala Ser	Ser	
1220			1225			1230			
Ala Thr	His Pro	Gly Thr	Gln	Thr Gly	Phe Thr	Val	Pro Ile	Arg	
1235			1240			1245			
Thr Val	Pro Ser	Ser Glu	Pro	Asp Thr	Met Ala	Ser	Trp Val	Thr	
1250			1255			1260			
His Pro	Pro Gln	Thr Ser	Thr	Pro Val	Ser Arg	Thr	Thr Ser	Ser	
1265			1270			1275			
Phe Ser	His Ser	Ser Pro	Asp	Ala Thr	Pro Val	Met	Ala Thr	Ser	
1280			1285			1290			
Pro Arg	Thr Glu	Ala Ser	Ser	Ala Val	Leu Thr	Thr	Ile Ser	Pro	
1295			1300			1305			
Gly Ala	Pro Glu	Met Val	Thr	Ser Gln	Ile Thr	Ser	Ser Gly	Ala	
1310			1315			1320			
Ala Thr	Ser Thr	Thr Val	Pro	Thr Leu	Thr His	Ser	Pro Gly	Met	
1325			1330			1335			
Pro Glu	Thr Thr	Ala Leu	Leu	Ser Thr	His Pro	Arg	Thr Glu	Thr	
1340			1345			1350			
Ser Lys	Thr Phe	Pro Ala	Ser	Thr Val	Phe Pro	Gln	Val Ser	Glu	

1355	1360	1365
Thr Thr Ala Ser Leu Thr	Ile Arg Pro Gly Ala	Glu Thr Ser Thr
1370	1375	1380
Ala Leu Pro Thr Gln Thr	Thr Ser Ser Leu Phe	Thr Leu Leu Val
1385	1390	1395
Thr Gly Thr Ser Arg Val	Asp Leu Ser Pro Thr	Ala Ser Pro Gly
1400	1405	1410
Val Ser Ala Lys Thr Ala	Pro Leu Ser Thr His	Pro Gly Thr Glu
1415	1420	1425
Thr Ser Thr Met Ile Pro	Thr Ser Thr Leu Ser	Leu Gly Leu Leu
1430	1435	1440
Glu Thr Thr Gly Leu Leu	Ala Thr Ser Ser Ser	Ala Glu Thr Ser
1445	1450	1455
Thr Ser Thr Leu Thr Leu	Thr Val Ser Pro Ala	Val Ser Gly Leu
1460	1465	1470
Ser Ser Ala Ser Ile Thr	Thr Asp Lys Pro Gln	Thr Val Thr Ser
1475	1480	1485
Trp Asn Thr Glu Thr Ser	Pro Ser Val Thr Ser	Val Gly Pro Pro
1490	1495	1500
Glu Phe Ser Arg Thr Val	Thr Gly Thr Thr Met	Thr Leu Ile Pro
1505	1510	1515
Ser Glu Met Pro Thr Pro	Pro Lys Thr Ser His	Gly Glu Gly Val
1520	1525	1530
Ser Pro Thr Thr Ile Leu	Arg Thr Thr Met Val	Glu Ala Thr Asn
1535	1540	1545
Leu Ala Thr Thr Gly Ser	Ser Pro Thr Val Ala	Lys Thr Thr Thr
1550	1555	1560
Thr Phe Asn Thr Leu Ala	Gly Ser Leu Phe Thr	Pro Leu Thr Thr
1565	1570	1575
Pro Gly Met Ser Thr Leu	Ala Ser Glu Ser Val	Thr Ser Arg Thr
1580	1585	1590
Ser Tyr Asn His Arg Ser	Trp Ile Ser Thr Thr	Ser Ser Tyr Asn
1595	1600	1605
Arg Arg Tyr Trp Thr Pro	Ala Thr Ser Thr Pro	Val Thr Ser Thr
1610	1615	1620
Phe Ser Pro Gly Ile Ser	Thr Ser Ser Ile Pro	Ser Ser Thr Ala
1625	1630	1635
Ala Thr Val Pro Phe Met	Val Pro Phe Thr Leu	Asn Phe Thr Ile
1640	1645	1650

Thr Asn	Leu Gln Tyr Glu	Glu Asp Met Arg His	Pro Gly Ser Arg	1655	1660	1665	
Lys Phe	Asn Ala Thr Glu Arg	Glu Leu Gln Gly Leu	Leu Lys Pro	1670	1675	1680	
Leu Phe	Arg Asn Ser Ser	Leu Glu Tyr Leu Tyr	Ser Gly Cys Arg	1685	1690	1695	
Leu Ala	Ser Leu Arg Pro	Glu Lys Asp Ser Ser	Ala Met Ala Val	1700	1705	1710	
Asp Ala	Ile Cys Thr His	Arg Pro Asp Pro Glu	Asp Leu Gly Leu	1715	1720	1725	
Asp Arg	Glu Arg Leu Tyr	Trp Glu Leu Ser Asn	Leu Thr Asn Gly	1730	1735	1740	
Ile Gln	Glu Leu Gly Pro	Tyr Thr Leu Asp Arg	Asn Ser Leu Tyr	1745	1750	1755	
Val Asn	Gly Phe Thr His	Arg Ser Ser Met Pro	Thr Thr Ser Thr	1760	1765	1770	
Pro Gly	Thr Ser Thr Val	Asp Val Gly Thr Ser	Gly Thr Pro Ser	1775	1780	1785	
Ser Ser	Pro Ser Pro Thr	Ala Ala Gly Pro Leu	Leu Met Pro Phe	1790	1795	1800	
Thr Leu	Asn Phe Thr Ile	Thr Asn Leu Gln Tyr	Glu Glu Asp Met	1805	1810	1815	
Arg Arg	Thr Gly Ser Arg	Lys Phe Asn Thr Met	Glu Ser Val Leu	1820	1825	1830	
Gln Gly	Leu Leu Lys Pro	Leu Phe Lys Asn Thr	Ser Val Gly Pro	1835	1840	1845	
Leu Tyr	Ser Gly Cys Arg	Leu Thr Leu Leu Arg	Pro Glu Lys Asp	1850	1855	1860	
Gly Ala	Ala Thr Gly Val	Asp Ala Ile Cys Thr	His Arg Leu Asp	1865	1870	1875	
Pro Lys	Ser Pro Gly Leu	Asn Arg Glu Gln Leu	Tyr Trp Glu Leu	1880	1885	1890	
Ser Lys	Leu Thr Asn Asp	Ile Glu Glu Leu Gly	Pro Tyr Thr Leu	1895	1900	1905	
Asp Arg	Asn Ser Leu Tyr	Val Asn Gly Phe Thr	His Gln Ser Ser	1910	1915	1920	
Val Ser	Thr Thr Ser Thr	Pro Gly Thr Ser Thr	Val Asp Leu Arg	1925	1930	1935	
Thr Ser	Gly Thr Pro Ser	Ser Leu Ser Ser Pro	Thr Ile Met Ala	1940	1945	1950	

Ala Gly 1955	Pro Leu Leu Val	Pro 1960	Phe Thr Leu Asn	Phe 1965	Thr Ile Thr
Asn Leu 1970	Gln Tyr Gly Glu	Asp 1975	Met Gly His Pro	Gly 1980	Ser Arg Lys
Phe Asn 1985	Thr Thr Glu Arg	Val 1990	Leu Gln Gly Leu	Leu 1995	Gly Pro Ile
Phe Lys 2000	Asn Thr Ser Val	Gly 2005	Pro Leu Tyr Ser	Gly 2010	Cys Arg Leu
Thr Ser 2015	Leu Arg Ser Glu	Lys 2020	Asp Gly Ala Ala	Thr 2025	Gly Val Asp
Ala Ile 2030	Cys Ile His His	Leu 2035	Asp Pro Lys Ser	Pro 2040	Gly Leu Asn
Arg Glu 2045	Arg Leu Tyr Trp	Glu 2050	Leu Ser Gln Leu	Thr 2055	Asn Gly Ile
Lys Glu 2060	Leu Gly Pro Tyr	Thr 2065	Leu Asp Arg Asn	Ser 2070	Leu Tyr Val
Asn Gly 2075	Phe Thr His Arg	Thr 2080	Ser Val Pro Thr	Ser 2085	Ser Thr Pro
Gly Thr 2090	Ser Thr Val Asp	Leu 2095	Gly Thr Ser Gly	Thr 2100	Pro Phe Ser
Leu Pro 2105	Ser Pro Ala Thr	Ala 2110	Gly Pro Leu Leu	Val 2115	Leu Phe Thr
Leu Asn 2120	Phe Thr Ile Thr	Asn 2125	Leu Lys Tyr Glu	Glu 2130	Asp Met His
Arg Pro 2135	Gly Ser Arg Lys	Phe 2140	Asn Thr Thr Glu	Arg 2145	Val Leu Gln
Thr Leu 2150	Leu Gly Pro Met	Phe 2155	Lys Asn Thr Ser	Val 2160	Gly Leu Leu
Tyr Ser 2165	Gly Cys Arg Leu	Thr 2170	Leu Leu Arg Ser	Glu 2175	Lys Asp Gly
Ala Ala 2180	Thr Gly Val Asp	Ala 2185	Ile Cys Thr His	Arg 2190	Leu Asp Pro
Lys Ser 2195	Pro Gly Leu Asp	Arg 2200	Glu Gln Leu Tyr	Trp 2205	Glu Leu Ser
Gln Leu 2210	Thr Asn Gly Ile	Lys 2215	Glu Leu Gly Pro	Tyr 2220	Thr Leu Asp
Arg Asn 2225	Ser Leu Tyr Val	Asn 2230	Gly Phe Thr His	Trp 2235	Ile Pro Val
Pro Thr	Ser Ser Thr Pro	Gly	Thr Ser Thr Val	Asp	Leu Gly Ser

2240					2245					2250				
Gly Thr	Pro	Ser	Ser	Leu	Pro	Ser	Pro	Thr	Ala	Ala	Gly	Pro	Leu	
2255					2260					2265				
Leu Val	Pro	Phe	Thr	Leu	Asn	Phe	Thr	Ile	Thr	Asn	Leu	Gln	Tyr	
2270					2275					2280				
Glu Glu	Asp	Met	His	His	Pro	Gly	Ser	Arg	Lys	Phe	Asn	Thr	Thr	
2285					2290					2295				
Glu Arg	Val	Leu	Gln	Gly	Leu	Leu	Gly	Pro	Met	Phe	Lys	Asn	Thr	
2300					2305					2310				
Ser Val	Gly	Leu	Leu	Tyr	Ser	Gly	Cys	Arg	Leu	Thr	Leu	Leu	Arg	
2315					2320					2325				
Ser Glu	Lys	Asp	Gly	Ala	Ala	Thr	Gly	Val	Asp	Ala	Ile	Cys	Thr	
2330					2335					2340				
His Arg	Leu	Asp	Pro	Lys	Ser	Pro	Gly	Val	Asp	Arg	Glu	Gln	Leu	
2345					2350					2355				
Tyr Trp	Glu	Leu	Ser	Gln	Leu	Thr	Asn	Gly	Ile	Lys	Glu	Leu	Gly	
2360					2365					2370				
Pro Tyr	Thr	Leu	Asp	Arg	Asn	Ser	Leu	Tyr	Val	Asn	Gly	Phe	Thr	
2375					2380					2385				
His Gln	Thr	Ser	Ala	Pro	Asn	Thr	Ser	Thr	Pro	Gly	Thr	Ser	Thr	
2390					2395					2400				
Val Asp	Leu	Gly	Thr	Ser	Gly	Thr	Pro	Ser	Ser	Leu	Pro	Ser	Pro	
2405					2410					2415				
Thr Ser	Ala	Gly	Pro	Leu	Leu	Val	Pro	Phe	Thr	Leu	Asn	Phe	Thr	
2420					2425					2430				
Ile Thr	Asn	Leu	Gln	Tyr	Glu	Glu	Asp	Met	Arg	His	Pro	Gly	Ser	
2435					2440					2445				
Arg Lys	Phe	Asn	Thr	Thr	Glu	Arg	Val	Leu	Gln	Gly	Leu	Leu	Lys	
2450					2455					2460				
Pro Leu	Phe	Lys	Ser	Thr	Ser	Val	Gly	Pro	Leu	Tyr	Ser	Gly	Cys	
2465					2470					2475				
Arg Leu	Thr	Leu	Leu	Arg	Ser	Glu	Lys	Asp	Gly	Ala	Ala	Thr	Gly	
2480					2485					2490				
Val Asp	Ala	Ile	Cys	Thr	His	Arg	Leu	Asp	Pro	Lys	Ser	Pro	Gly	
2495					2500					2505				
Val Asp	Arg	Glu	Gln	Leu	Tyr	Trp	Glu	Leu	Ser	Gln	Leu	Thr	Asn	
2510					2515					2520				
Gly Ile	Lys	Glu	Leu	Gly	Pro	Tyr	Thr	Leu	Asp	Arg	Asn	Ser	Leu	
2525					2530					2535				

Tyr Val Asn Gly Phe Thr His Gln Thr Ser Ala Pro Asn Thr Ser
 2540 2545 2550
 Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro
 2555 2560 2565
 Ser Ser Leu Pro Ser Pro Thr Ser Ala Gly Pro Leu Leu Val Pro
 2570 2575 2580
 Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp
 2585 2590 2595
 Met His His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val
 2600 2605 2610
 Leu Gln Gly Leu Leu Gly Pro Met Phe Lys Asn Thr Ser Val Gly
 2615 2620 2625
 Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys
 2630 2635 2640
 Asn Gly Ala Ala Thr Gly Met Asp Ala Ile Cys Ser His Arg Leu
 2645 2650 2655
 Asp Pro Lys Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp Glu
 2660 2665 2670
 Leu Ser Gln Leu Thr His Gly Ile Lys Glu Leu Gly Pro Tyr Thr
 2675 2680 2685
 Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser
 2690 2695 2700
 Ser Val Ala Pro Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu
 2705 2710 2715
 Gly Thr Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr Thr Ala
 2720 2725 2730
 Val Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn
 2735 2740 2745
 Leu Gln Tyr Gly Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe
 2750 2755 2760
 Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Leu Phe
 2765 2770 2775
 Lys Asn Ser Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Ile
 2780 2785 2790
 Ser Leu Arg Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala
 2795 2800 2805
 Ile Cys Thr His His Leu Asn Pro Gln Ser Pro Gly Leu Asp Arg
 2810 2815 2820
 Glu Gln Leu Tyr Trp Gln Leu Ser Gln Met Thr Asn Gly Ile Lys
 2825 2830 2835

Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn
 2840 2845 2850
 Gly Phe Thr His Arg Ser Ser Gly Leu Thr Thr Ser Thr Pro Trp
 2855 2860 2865
 Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Pro Val
 2870 2875 2880
 Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Val Pro Phe Thr Leu
 2885 2890 2895
 Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met His Arg
 2900 2905 2910
 Pro Gly Ser Arg Lys Phe Asn Ala Thr Glu Arg Val Leu Gln Gly
 2915 2920
 Leu Leu Ser Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu Tyr
 2930 2935 2940
 Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu Lys Asp Gly Ala
 2945 2950 2955
 Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro Asn Pro Lys
 2960 2965 2970
 Arg Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln
 2975 2980 2985
 Leu Thr His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp Arg
 2990 2995 3000
 Asp Ser Leu Tyr Val Asn Gly Phe Thr His Gln Asn Ser Val Pro
 3005 3010 3015
 Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr
 3020 3025 3030
 Gly Thr Pro Ser Ser Phe Pro Gly His Thr Glu Pro Gly Pro Leu
 3035 3040 3045
 Leu Ile Pro Phe Thr Phe Asn Phe Thr Ile Thr Asn Leu His Tyr
 3050 3055 3060
 Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr
 3065 3070 3075
 Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr
 3080 3085 3090
 Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg
 3095 3100 3105
 Pro Glu Lys Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu
 3110 3115 3120
 Tyr His Pro Asn Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu

3125	3130	3135
Tyr Cys Glu Leu Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly		
3140	3145	3150
Pro Tyr Ser Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr		
3155	3160	3165
His Gln Asn Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr		
3170	3175	3180
Val Tyr Trp Ala Thr Thr Gly Thr Pro Ser Ser Phe Pro Gly His		
3185	3190	3195
Thr Glu Pro Gly Pro Leu Leu Ile Pro Phe Thr Phe Asn Phe Thr		
3200	3205	3210
Ile Thr Asn Leu His Tyr Glu Glu Asn Met Gln His Pro Gly Ser		
3215	3220	3225
Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Lys		
3230	3235	3240
Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys		
3245	3250	3255
Arg Leu Thr Leu Leu Arg Pro Glu Lys His Glu Ala Ala Thr Gly		
3260	3265	3270
Val Asp Thr Ile Cys Thr His Arg Val Asp Pro Ile Gly Pro Gly		
3275	3280	3285
Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn		
3290	3295	3300
Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu		
3305	3310	3315
Tyr Val Asn Gly Phe Asn Pro Arg Ser Ser Val Pro Thr Thr Ser		
3320	3325	3330
Thr Pro Gly Thr Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro		
3335	3340	3345
Ser Ser Leu Pro Gly His Thr Ala Pro Val Pro Leu Leu Ile Pro		
3350	3355	3360
Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu His Tyr Glu Glu Asn		
3365	3370	3375
Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val		
3380	3385	3390
Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Gly		
3395	3400	3405
Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys		
3410	3415	3420

His Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Val
 3425 3430 3435
 Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Xaa Leu Tyr Trp Glu
 3440 3445 3450
 Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa
 3455 3460 3465
 Leu Asp Arg Xaa Ser Leu Tyr Val Asn Gly Phe Xaa Xaa Xaa Xaa
 3470 3475 3480
 Xaa Xaa Xaa Xaa Thr Ser Thr Pro Gly Thr Ser Xaa Val Xaa Leu
 3485 3490 3495
 Xaa Thr Ser Gly Thr Pro Xaa Xaa Xaa Pro Xaa Xaa Thr Ser Ala
 3500 3505 3510
 Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn
 3515 3520 3525
 Leu Gln Tyr Glu Glu Asp Met His His Pro Gly Ser Arg Lys Phe
 3530 3535 3540
 Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Met Phe
 3545 3550 3555
 Lys Asn Thr Ser Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr
 3560 3565 3570
 Leu Leu Arg Pro Glu Lys Asn Gly Ala Ala Thr Gly Met Asp Ala
 3575 3580 3585
 Ile Cys Ser His Arg Leu Asp Pro Lys Ser Pro Gly Leu Asp Arg
 3590 3595 3600
 Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His Gly Ile Lys
 3605 3610 3615
 Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn
 3620 3625 3630
 Gly Phe Thr His Arg Ser Ser Val Ala Pro Thr Ser Thr Pro Gly
 3635 3640 3645
 Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Ser Leu
 3650 3655 3660
 Pro Ser Pro Thr Thr Ala Val Pro Leu Leu Val Pro Phe Thr Leu
 3665 3670 3675
 Asn Phe Thr Ile Thr Asn Leu Gln Tyr Gly Glu Asp Met Arg His
 3680 3685 3690
 Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly
 3695 3700 3705
 Leu Leu Gly Pro Leu Phe Lys Asn Ser Ser Val Gly Pro Leu Tyr
 3710 3715 3720

Ser Gly Cys Arg Leu Ile Ser Leu Arg Ser Glu Lys Asp Gly Ala
 3725 3730 3735
 Ala Thr Gly Val Asp Ala Ile Cys Thr His His Leu Asn Pro Gln
 3740 3745 3750
 Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Gln Leu Ser Gln
 3755 3760 3765
 Met Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg
 3770 3775 3780
 Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Gly Leu
 3785 3790 3795
 Thr Thr Ser Thr Pro Trp Thr Ser Thr Val Asp Leu Gly Thr Ser
 3800 3805 3810
 Gly Thr Pro Ser Pro Val Pro Ser Pro Thr Thr Ala Gly Pro Leu
 3815 3820 3825
 Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr
 3830 3835 3840
 Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe Asn Ala Thr
 3845 3850 3855
 Glu Arg Val Leu Gln Gly Leu Leu Ser Pro Ile Phe Lys Asn Ser
 3860 3865 3870
 Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg
 3875 3880 3885
 Pro Glu Lys Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu
 3890 3895 3900
 Tyr His Pro Asn Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu
 3905 3910 3915
 Tyr Trp Glu Leu Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly
 3920 3925 3930
 Pro Tyr Ser Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr
 3935 3940 3945
 His Gln Ser Ser Met Thr Thr Thr Arg Thr Pro Asp Thr Ser Thr
 3950 3955 3960
 Met His Leu Ala Thr Ser Arg Thr Pro Ala Ser Leu Ser Gly Pro
 3965 3970 3975
 Thr Thr Ala Ser Pro Leu Leu Val Leu Phe Thr Ile Asn Cys Thr
 3980 3985 3990
 Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr Gly Ser
 3995 4000 4005
 Arg Lys Phe Asn Thr Met Glu Ser Val Leu Gln Gly Leu Leu Lys

4010	4015	4020
Pro Leu Phe Lys Asn Thr Ser	Val Gly Pro Leu Tyr	Ser Gly Cys
4025	4030	4035
Arg Leu Thr Leu Leu Arg Pro	Lys Lys Asp Gly Ala	Ala Thr Gly
4040	4045	4050
Val Asp Ala Ile Cys Thr His	Arg Leu Asp Pro Lys	Ser Pro Gly
4055	4060	4065
Leu Asn Arg Glu Gln Leu Tyr	Trp Glu Leu Ser Lys	Leu Thr Asn
4070	4075	4080
Asp Ile Glu Glu Leu Gly Pro	Tyr Thr Leu Asp Arg	Asn Ser Leu
4085	4090	4095
Tyr Val Asn Gly Phe Thr His	Gln Ser Ser Val Ser	Thr Thr Ser
4100	4105	4110
Thr Pro Gly Thr Ser Thr Val	Asp Leu Arg Thr Ser	Gly Thr Pro
4115	4120	4125
Ser Ser Leu Ser Ser Pro Thr	Ile Met Xaa Xaa Xaa	Pro Leu Leu
4130	4135	4140
Xaa Pro Phe Thr Leu Asn Phe	Thr Ile Thr Asn Leu	Xaa Tyr Glu
4145	4150	4155
Glu Xaa Met Xaa Xaa Pro Gly	Ser Arg Lys Phe Asn	Thr Thr Glu
4160	4165	4170
Arg Val Leu Gln Gly Leu Leu	Arg Pro Leu Phe Lys	Asn Thr Ser
4175	4180	4185
Val Ser Ser Leu Tyr Ser Gly	Cys Arg Leu Thr Leu	Leu Arg Pro
4190	4195	4200
Glu Lys Asp Gly Ala Ala Thr	Arg Val Asp Ala Ala	Cys Thr Tyr
4205	4210	4215
Arg Pro Asp Pro Lys Ser Pro	Gly Leu Asp Arg Glu	Gln Leu Tyr
4220	4225	4230
Trp Glu Leu Ser Gln Leu Thr	His Ser Ile Thr Glu	Leu Gly Pro
4235	4240	4245
Tyr Thr Leu Asp Arg Val Ser	Leu Tyr Val Asn Gly	Phe Asn Pro
4250	4255	4260
Arg Ser Ser Val Pro Thr Thr	Ser Thr Pro Gly Thr	Ser Thr Val
4265	4270	4275
His Leu Ala Thr Ser Gly Thr	Pro Ser Ser Leu Pro	Gly His Thr
4280	4285	4290
Xaa Xaa Xaa Pro Leu Leu Xaa	Pro Phe Thr Leu Asn	Phe Thr Ile
4295	4300	4305

Thr	Asn	Leu	Xaa	Tyr	Glu	Glu	Xaa	Met	Xaa	Xaa	Pro	Gly	Ser	Arg	
4310						4315					4320				
Lys	Phe	Asn	Thr	Thr	Glu	Arg	Val	Leu	Gln	Gly	Leu	Leu	Lys	Pro	
4325						4330					4335				
Leu	Phe	Arg	Asn	Ser	Ser	Leu	Glu	Tyr	Leu	Tyr	Ser	Gly	Cys	Arg	
4340						4345					4350				
Leu	Ala	Ser	Leu	Arg	Pro	Glu	Lys	Asp	Ser	Ser	Ala	Met	Ala	Val	
4355						4360					4365				
Asp	Ala	Ile	Cys	Thr	His	Arg	Pro	Asp	Pro	Glu	Asp	Leu	Gly	Leu	
4370						4375					4380				
Asp	Arg	Glu	Arg	Leu	Tyr	Trp	Glu	Leu	Ser	Asn	Leu	Thr	Asn	Gly	
4385						4390					4395				
Ile	Gln	Glu	Leu	Gly	Pro	Tyr	Thr	Leu	Asp	Arg	Asn	Ser	Leu	Tyr	
4400						4405					4410				
Val	Asn	Gly	Phe	Thr	His	Arg	Ser	Ser	Phe	Leu	Thr	Thr	Ser	Thr	
4415						4420					4425				
Pro	Trp	Thr	Ser	Thr	Val	Asp	Leu	Gly	Thr	Ser	Gly	Thr	Pro	Ser	
4430						4435					4440				
Pro	Val	Pro	Ser	Pro	Thr	Thr	Ala	Gly	Pro	Leu	Leu	Val	Pro	Phe	
4445						4450					4455				
Thr	Leu	Asn	Phe	Thr	Ile	Thr	Asn	Leu	Gln	Tyr	Glu	Glu	Asp	Met	
4460						4465					4470				
His	Arg	Pro	Gly	Ser	Arg	Arg	Phe	Asn	Thr	Thr	Glu	Arg	Val	Leu	
4475						4480					4485				
Gln	Gly	Leu	Leu	Thr	Pro	Leu	Phe	Lys	Asn	Thr	Ser	Val	Gly	Pro	
4490						4495					4500				
Leu	Tyr	Ser	Gly	Cys	Arg	Leu	Thr	Leu	Leu	Arg	Pro	Glu	Lys	Gln	
4505						4510					4515				
Glu	Ala	Ala	Thr	Gly	Val	Asp	Thr	Ile	Cys	Thr	His	Arg	Val	Asp	
4520						4525					4530				
Pro	Ile	Gly	Pro	Gly	Leu	Asp	Arg	Glu	Arg	Leu	Tyr	Trp	Glu	Leu	
4535						4540					4545				
Ser	Gln	Leu	Thr	Asn	Ser	Ile	Thr	Glu	Leu	Gly	Pro	Tyr	Thr	Leu	
4550						4555					4560				
Asp	Arg	Asp	Ser	Leu	Tyr	Val	Asn	Gly	Phe	Asn	Pro	Trp	Ser	Ser	
4565						4570					4575				
Val	Pro	Thr	Thr	Ser	Thr	Pro	Gly	Thr	Ser	Thr	Val	His	Leu	Ala	
4580						4585					4590				
Thr	Ser	Gly	Thr	Pro	Ser	Ser	Leu	Pro	Gly	His	Thr	Ala	Pro	Val	
4595						4600					4605				

Pro Leu	Leu Ile	Pro Phe	Thr	Leu Asn	Phe Thr	Ile	Thr Asp	Leu	
4610			4615			4620			
His Tyr	Glu Glu	Asn Met	Gln	His Pro	Gly Ser	Arg	Lys Phe	Asn	
4625			4630			4635			
Thr Thr	Glu Arg	Val Leu	Gln	Gly Leu	Leu Lys	Pro	Leu Phe	Lys	
4640			4645			4650			
Ser Thr	Ser Val	Gly Pro	Leu	Tyr Ser	Gly Cys	Arg	Leu Thr	Leu	
4655			4660			4665			
Leu Arg	Pro Glu	Lys His	Gly	Ala Ala	Thr Gly	Val	Asp Ala	Ile	
4670			4675			4680			
Cys Thr	Leu Arg	Leu Asp	Pro	Thr Gly	Pro Gly	Leu	Asp Arg	Glu	
4685			4690			4695			
Arg Leu	Tyr Trp	Glu Leu	Ser	Gln Leu	Thr Asn	Ser	Val Thr	Glu	
4700			4705			4710			
Leu Gly	Pro Tyr	Thr Leu	Asp	Arg Asp	Ser Leu	Tyr	Val Asn	Gly	
4715			4720			4725			
Phe Thr	His Arg	Ser Ser	Val	Pro Thr	Thr Ser	Ile	Pro Gly	Thr	
4730			4735			4740			
Ser Ala	Val His	Leu Glu	Thr	Ser Gly	Thr Pro	Ala	Ser Leu	Pro	
4745			4750			4755			
Gly His	Thr Ala	Pro Gly	Pro	Leu Leu	Val Pro	Phe	Thr Leu	Asn	
4760			4765			4770			
Phe Thr	Ile Thr	Asn Leu	Gln	Tyr Glu	Glu Asp	Met	Arg His	Pro	
4775			4780			4785			
Gly Ser	Arg Lys	Phe Ser	Thr	Thr Glu	Arg Val	Leu	Gln Gly	Leu	
4790			4795			4800			
Leu Lys	Pro Leu	Phe Lys	Asn	Thr Ser	Val Ser	Ser	Leu Tyr	Ser	
4805			4810			4815			
Gly Cys	Arg Leu	Thr Leu	Leu	Arg Pro	Glu Lys	Asp	Gly Ala	Ala	
4820			4825			4830			
Thr Arg	Val Asp	Ala Val	Cys	Thr His	Arg Pro	Asp	Pro Lys	Ser	
4835			4840			4845			
Pro Gly	Leu Asp	Arg Glu	Arg	Leu Tyr	Trp Lys	Leu	Ser Gln	Leu	
4850			4855			4860			
Thr His	Gly Ile	Thr Glu	Leu	Gly Pro	Tyr Thr	Leu	Asp Arg	His	
4865			4870			4875			
Ser Leu	Tyr Val	Asn Gly	Phe	Thr His	Gln Ser	Ser	Met Thr	Thr	
4880			4885			4890			
Thr Arg	Thr Pro	Asp Thr	Ser	Thr Met	His Leu	Ala	Thr Ser	Arg	

4895	4900	4905
Thr Pro Ala Ser Leu Ser Gly 4910	Pro Thr Thr Ala Ser 4915	Pro Leu Leu 4920
Val Leu Phe Thr Ile Asn Phe 4925	Thr Ile Thr Asn Gln 4930	Arg Tyr Glu 4935
Glu Asn Met His His Pro Gly 4940	Ser Arg Lys Phe Asn 4945	Thr Thr Glu 4950
Arg Val Leu Gln Gly Leu Leu 4955	Arg Pro Val Phe Lys 4960	Asn Thr Ser 4965
Val Gly Pro Leu Tyr Ser Gly 4970	Cys Arg Leu Thr Leu 4975	Leu Arg Pro 4980
Lys Lys Asp Gly Ala Ala Thr 4985	Lys Val Asp Ala Ile 4990	Cys Thr Tyr 4995
Arg Pro Asp Pro Lys Ser Pro 5000	Gly Leu Asp Arg Glu 5005	Gln Leu Tyr 5010
Trp Glu Leu Ser Gln Leu Thr 5015	His Ser Ile Thr Glu 5020	Leu Gly Pro 5025
Tyr Thr Gln Asp Arg Asp Ser 5030	Leu Tyr Val Asn Gly 5035	Phe Thr His 5040
Arg Ser Ser Val Pro Thr Thr 5045	Ser Ile Pro Gly Thr 5050	Ser Ala Val 5055
His Leu Glu Thr Ser Gly Thr 5060	Pro Ala Ser Leu Pro 5065	Gly His Thr 5070
Ala Pro Gly Pro Leu Leu Val 5075	Pro Phe Thr Leu Asn 5080	Phe Thr Ile 5085
Thr Asn Leu Gln Tyr Glu Glu 5090	Asp Met Arg His Pro 5095	Gly Ser Arg 5100
Lys Phe Asn Thr Thr Glu Arg 5105	Val Leu Gln Gly Leu 5110	Leu Lys Pro 5115
Leu Phe Lys Ser Thr Ser Val 5120	Gly Pro Leu Tyr Ser 5125	Gly Cys Arg 5130
Leu Thr Leu Leu Arg Pro Glu 5135	Lys Arg Gly Ala Ala 5140	Thr Gly Val 5145
Asp Thr Ile Cys Thr His Arg 5150	Leu Asp Pro Leu Asn 5155	Pro Gly Leu 5160
Asp Arg Glu Gln Leu Tyr Trp 5165	Glu Leu Ser Lys Leu 5170	Thr Arg Gly 5175
Ile Ile Glu Leu Gly Pro Tyr 5180	Leu Leu Asp Arg Gly 5185	Ser Leu Tyr 5190

Val Asn Gly Phe Thr His Arg Thr Ser Val Pro Thr Thr Ser Thr
 5195 5200 5205
 Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Phe
 5210 5215 5220
 Ser Leu Pro Ser Pro Ala Xaa Xaa Xaa Pro Leu Leu Xaa Pro Phe
 5225 5230 5235
 Thr Leu Asn Phe Thr Ile Thr Asn Leu Xaa Tyr Glu Glu Xaa Met
 5240 5245 5250
 Xaa Xaa Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu
 5255 5260 5265
 Gln Thr Leu Leu Gly Pro Met Phe Lys Asn Thr Ser Val Gly Leu
 5270 5275 5280
 Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Ser Glu Lys Asp
 5285 5290 5295
 Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu Asp
 5300 5305 5310
 Pro Lys Ser Pro Gly Val Asp Arg Glu Gln Leu Tyr Trp Glu Leu
 5315 5320 5325
 Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu
 5330 5335 5340
 Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Trp Ile Pro
 5345 5350 5355
 Val Pro Thr Ser Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly
 5360 5365 5370
 Ser Gly Thr Pro Ser Leu Pro Ser Ser Pro Thr Thr Ala Gly Pro
 5375 5380 5385
 Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Lys
 5390 5395 5400
 Tyr Glu Glu Asp Met His Cys Pro Gly Ser Arg Lys Phe Asn Thr
 5405 5410 5415
 Thr Glu Arg Val Leu Gln Ser Leu Leu Gly Pro Met Phe Lys Asn
 5420 5425 5430
 Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu
 5435 5440 5445
 Arg Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys
 5450 5455 5460
 Thr His Arg Leu Asp Pro Lys Ser Pro Gly Val Asp Arg Glu Gln
 5465 5470 5475
 Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu
 5480 5485 5490

Gly	Pro	Tyr	Thr	Leu	Asp	Arg	Asn	Ser	Leu	Tyr	Val	Asn	Gly	Phe
5495						5500					5505			
Thr	His	Gln	Thr	Ser	Ala	Pro	Asn	Thr	Ser	Thr	Pro	Gly	Thr	Ser
5510						5515					5520			
Thr	Val	Asp	Leu	Gly	Thr	Ser	Gly	Thr	Pro	Ser	Ser	Leu	Pro	Ser
5525						5530					5535			
Pro	Thr	Xaa	Xaa	Xaa	Pro	Leu	Leu	Xaa	Pro	Phe	Thr	Leu	Asn	Phe
5540						5545					5550			
Thr	Ile	Thr	Asn	Leu	Xaa	Tyr	Glu	Glu	Xaa	Met	Xaa	Xaa	Pro	Gly
5555						5560					5565			
Ser	Arg	Lys	Phe	Asn	Thr	Thr	Glu	Arg	Val	Leu	Gln	Gly	Leu	Leu
5570						5575					5580			
Xaa	Pro	Xaa	Phe	Lys	Xaa	Thr	Ser	Val	Gly	Xaa	Leu	Tyr	Ser	Gly
5585						5590					5595			
Cys	Arg	Leu	Thr	Leu	Leu	Arg	Xaa	Glu	Lys	Xaa	Xaa	Ala	Ala	Thr
5600						5605					5610			
Xaa	Val	Asp	Xaa	Xaa	Cys	Xaa	Xaa	Xaa	Xaa	Asp	Pro	Xaa	Xaa	Pro
5615						5620					5625			
Gly	Leu	Asp	Arg	Glu	Xaa	Leu	Tyr	Trp	Glu	Leu	Ser	Xaa	Leu	Thr
5630						5635					5640			
Xaa	Xaa	Ile	Xaa	Glu	Leu	Gly	Pro	Tyr	Xaa	Leu	Asp	Arg	Xaa	Ser
5645						5650					5655			
Leu	Tyr	Val	Asn	Gly	Phe	Thr	His	Trp	Ile	Pro	Val	Pro	Thr	Ser
5660						5665					5670			
Ser	Thr	Pro	Gly	Thr	Ser	Thr	Val	Asp	Leu	Gly	Ser	Gly	Thr	Pro
5675						5680					5685			
Ser	Ser	Leu	Pro	Ser	Pro	Thr	Thr	Ala	Gly	Pro	Leu	Leu	Val	Pro
5690						5695					5700			
Phe	Thr	Leu	Asn	Phe	Thr	Ile	Thr	Asn	Leu	Lys	Tyr	Glu	Glu	Asp
5705						5710					5715			
Met	His	Cys	Pro	Gly	Ser	Arg	Lys	Phe	Asn	Thr	Thr	Glu	Arg	Val
5720						5725					5730			
Leu	Gln	Ser	Leu	Leu	Gly	Pro	Met	Phe	Lys	Asn	Thr	Ser	Val	Gly
5735						5740					5745			
Pro	Leu	Tyr	Ser	Gly	Cys	Arg	Leu	Thr	Ser	Leu	Arg	Ser	Glu	Lys
5750						5755					5760			
Asp	Gly	Ala	Ala	Thr	Gly	Val	Asp	Ala	Ile	Cys	Thr	His	Arg	Val
5765						5770					5775			
Asp	Pro	Lys	Ser	Pro	Gly	Val	Asp	Arg	Glu	Gln	Leu	Tyr	Trp	Gln
5785						5790					5795			

5780	5785	5790
Leu Ser Gln Leu Thr Asn Gly	Ile Lys Glu Leu Gly	Pro Tyr Thr
5795	5800	5805
Leu Asp Arg Asn Ser Leu Tyr	Val Asn Gly Phe Thr	His Gln Thr
5810	5815	5820
Ser Ala Pro Asn Thr Ser Thr	Pro Gly Thr Ser Thr	Val Asp Leu
5825	5830	5835
Gly Thr Ser Gly Thr Pro Ser	Ser Leu Pro Ser Pro	Thr Ser Ala
5840	5845	5850
Gly Pro Leu Leu Val Pro Phe	Thr Leu Asn Phe Thr	Ile Thr Asn
5855	5860	5865
Leu Gln Tyr Glu Glu Asp Met	His His Pro Gly Ser	Arg Lys Phe
5870	5875	5880
Asn Thr Thr Glu Arg Val Leu	Gln Gly Leu Leu Gly	Pro Met Phe
5885	5890	5895
Lys Asn Thr Ser Val Gly Leu	Leu Tyr Ser Gly Cys	Arg Leu Thr
5900	5905	5910
Leu Leu Arg Pro Glu Lys Asn	Gly Ala Ala Thr Gly	Met Asp Ala
5915	5920	5925
Ile Cys Thr His Arg Leu Asp	Pro Lys Ser Pro Gly	Leu Asp Arg
5930	5935	5940
Glu Xaa Leu Tyr Trp Glu Leu	Ser Xaa Leu Thr Xaa	Xaa Ile Xaa
5945	5950	5955
Glu Leu Gly Pro Tyr Xaa Leu	Asp Arg Xaa Ser Leu	Tyr Val Asn
5960	5965	5970
Gly Phe Xaa Xaa Xaa Xaa Xaa	Xaa Xaa Xaa Thr Ser	Thr Pro Gly
5975	5980	5985
Thr Ser Xaa Val Xaa Leu Xaa	Thr Ser Gly Thr Pro	Xaa Xaa Xaa
5990	5995	6000
Pro Xaa Xaa Thr Xaa Xaa Xaa	Pro Leu Leu Xaa Pro	Phe Thr Leu
6005	6010	6015
Asn Phe Thr Ile Thr Asn Leu	Xaa Tyr Glu Glu Xaa	Met Xaa Xaa
6020	6025	6030
Pro Gly Ser Arg Lys Phe Asn	Thr Thr Glu Arg Val	Leu Gln Gly
6035	6040	6045
Leu Leu Lys Pro Leu Phe Arg	Asn Ser Ser Leu Glu	Tyr Leu Tyr
6050	6055	6060
Ser Gly Cys Arg Leu Ala Ser	Leu Arg Pro Glu Lys	Asp Ser Ser
6065	6070	6075

Ala Met	Ala Val	Asp Ala	Ile Cys	Thr His	Arg Pro	Asp Pro	Glu
6080			6085		6090		
Asp Leu	Gly Leu	Asp Arg	Glu Arg	Leu Tyr	Trp Glu	Leu Ser	Asn
6095			6100		6105		
Leu Thr	Asn Gly	Ile Gln	Glu Leu	Gly Pro	Tyr Thr	Leu Asp	Arg
6110			6115		6120		
Asn Ser	Leu Tyr	Val Asn	Gly Phe	Thr His	Arg Ser	Ser Met	Pro
6125			6130		6135		
Thr Thr	Ser Thr	Pro Gly	Thr Ser	Thr Val	Asp Val	Gly Thr	Ser
6140			6145		6150		
Gly Thr	Pro Ser	Ser Ser	Pro Ser	Pro Thr	Thr Ala	Gly Pro	Leu
6155			6160		6165		
Leu Ile	Pro Phe	Thr Leu	Asn Phe	Thr Ile	Thr Asn	Leu Gln	Tyr
6170			6175		6180		
Gly Glu	Asp Met	Gly His	Pro Gly	Ser Arg	Lys Phe	Asn Thr	Thr
6185			6190		6195		
Glu Arg	Val Leu	Gln Gly	Leu Leu	Gly Pro	Ile Phe	Lys Asn	Thr
6200			6205		6210		
Ser Val	Gly Pro	Leu Tyr	Ser Gly	Cys Arg	Leu Thr	Ser Leu	Arg
6215			6220		6225		
Ser Glu	Lys Asp	Gly Ala	Ala Thr	Gly Val	Asp Ala	Ile Cys	Ile
6230			6235		6240		
His His	Leu Asp	Pro Lys	Ser Pro	Gly Leu	Asn Arg	Glu Arg	Leu
6245			6250		6255		
Tyr Trp	Glu Leu	Ser Gln	Leu Thr	Asn Gly	Ile Lys	Glu Leu	Gly
6260			6265		6270		
Pro Tyr	Thr Leu	Asp Arg	Asn Ser	Leu Tyr	Val Asn	Gly Phe	Thr
6275			6280		6285		
His Arg	Thr Ser	Val Pro	Thr Thr	Ser Thr	Pro Gly	Thr Ser	Thr
6290			6295		6300		
Val Asp	Leu Gly	Thr Ser	Gly Thr	Pro Phe	Ser Leu	Pro Ser	Pro
6305			6310		6315		
Ala Thr	Ala Gly	Pro Leu	Leu Val	Leu Phe	Thr Leu	Asn Phe	Thr
6320			6325		6330		
Ile Thr	Asn Leu	Lys Tyr	Glu Glu	Asp Met	His Arg	Pro Gly	Ser
6335			6340		6345		
Arg Lys	Phe Asn	Thr Thr	Glu Arg	Val Leu	Gln Thr	Leu Leu	Gly
6350			6355		6360		
Pro Met	Phe Lys	Asn Thr	Ser Val	Gly Leu	Leu Tyr	Ser Gly	Cys
6365			6370		6375		

Arg Leu Thr Leu Leu Arg Ser Glu Lys Asp Gly Ala Ala Thr Gly
 6380 6385 6390
 Val Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser Pro Gly
 6395 6400 6405
 Leu Asp Arg Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa
 6410 6415 6420
 Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa Leu Asp Arg Xaa Ser Leu
 6425 6430 6435
 Tyr Val Asn Gly Phe Xaa Xaa Xaa Xaa Xaa Xaa Thr Ser
 6440 6445 6450
 Thr Pro Gly Thr Ser Xaa Val Xaa Leu Xaa Thr Ser Gly Thr Pro
 6455 6460 6465
 Xaa Xaa Xaa Pro Xaa Xaa Thr Xaa Xaa Xaa Pro Leu Leu Xaa Pro
 6470 6475 6480
 Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Xaa Tyr Glu Glu Xaa
 6485 6490 6495
 Met Xaa Xaa Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val
 6500 6505 6510
 Leu Gln Gly Leu Leu Arg Pro Val Phe Lys Asn Thr Ser Val Gly
 6515 6520 6525
 Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys Lys
 6530 6535 6540
 Asp Gly Ala Ala Thr Lys Val Asp Ala Ile Cys Thr Tyr Arg Pro
 6545 6550 6555
 Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu
 6560 6565 6570
 Leu Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr
 6575 6580 6585
 Gln Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser
 6590 6595 6600
 Ser Val Pro Thr Thr Ser Ile Pro Gly Thr Ser Ala Val His Leu
 6605 6610 6615
 Glu Thr Thr Gly Thr Pro Ser Ser Phe Pro Gly His Thr Glu Pro
 6620 6625 6630
 Gly Pro Leu Leu Ile Pro Phe Thr Phe Asn Phe Thr Ile Thr Asn
 6635 6640 6645
 Leu Arg Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe
 6650 6655 6660
 Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Thr Pro Leu Phe

6665	6670	6675
Lys Asn Thr Ser Val Gly Pro 6680	Leu Tyr Ser Gly Cys Arg Leu Thr 6685	
Leu Leu Arg Pro Glu Lys Gln 6695	Glu Ala Ala Thr Gly Val Asp Thr 6700	
Ile Cys Thr His Arg Val Asp 6710	Pro Ile Gly Pro Gly Leu Asp Arg 6715	
Glu Arg Leu Tyr Trp Glu 6725	Leu Ser Gln Leu Thr Asn Ser Ile Thr 6730	
Glu Leu Gly Pro Tyr Thr 6740	Leu Asp Arg Asp Ser Leu Tyr Val Asp 6745	
Gly Phe Asn Pro Trp Ser 6755	Ser Val Pro Thr Thr Ser Thr Pro Gly 6760	
Thr Ser Thr Val His Leu 6770	Ala Thr Ser Gly Thr Pro Ser Pro Leu 6775	
Pro Gly His Thr Ala Pro 6785	Val Pro Leu Leu Ile Pro Phe Thr Leu 6790	
Asn Phe Thr Ile Thr Asp 6800	Leu His Tyr Glu Glu Asn Met Gln His 6805	
Pro Gly Ser Arg Lys Phe 6815	Asn Thr Thr Glu Arg Val Leu Gln Gly 6820	
Leu Leu Lys Pro Leu Phe 6830	Lys Ser Thr Ser Val Gly Pro Leu Tyr 6835	
Ser Gly Cys Arg Leu Thr 6845	Leu Leu Arg Pro Glu Lys His Gly Ala 6850	
Ala Thr Gly Val Asp Ala 6860	Ile Cys Thr Leu Arg Leu Asp Pro Thr 6865	
Gly Pro Gly Leu Asp Arg 6875	Glu Arg Leu Tyr Trp Glu Leu Ser Gln 6880	
Leu Thr Asn Ser Ile Thr 6890	Glu Leu Gly Pro Tyr Thr Leu Asp Arg 6895	
Asp Ser Leu Tyr Val Asn 6905	Gly Phe Asn Pro Trp Ser Ser Val Pro 6910	
Thr Thr Ser Thr Pro Gly 6920	Thr Ser Thr Val His Leu Ala Thr Ser 6925	
Gly Thr Pro Ser Ser Leu 6935	Pro Gly His Thr Thr Ala Gly Pro Leu 6940	
Leu Val Pro Phe Thr Leu 6950	Asn Phe Thr Ile Thr Asn Leu Lys Tyr 6955	

Glu	Glu	Asp	Met	His	Cys	Pro	Gly	Ser	Arg	Lys	Phe	Asn	Thr	Thr
6965						6970					6975			
Glu	Arg	Val	Leu	Gln	Ser	Leu	His	Gly	Pro	Met	Phe	Lys	Asn	Thr
6980						6985					6990			
Ser	Val	Gly	Pro	Leu	Tyr	Ser	Gly	Cys	Arg	Leu	Thr	Leu	Leu	Arg
6995						7000					7005			
Ser	Glu	Lys	Asp	Gly	Ala	Ala	Thr	Gly	Val	Asp	Ala	Ile	Cys	Thr
7010						7015					7020			
His	Arg	Leu	Asp	Pro	Lys	Ser	Pro	Gly	Leu	Asp	Arg	Glu	Xaa	Leu
7025						7030					7035			
Tyr	Trp	Glu	Leu	Ser	Xaa	Leu	Thr	Xaa	Xaa	Ile	Xaa	Glu	Leu	Gly
7040						7045					7050			
Pro	Tyr	Xaa	Leu	Asp	Arg	Xaa	Ser	Leu	Tyr	Val	Asn	Gly	Phe	Xaa
7055						7060					7065			
Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Thr	Ser	Thr	Pro	Gly	Thr	Ser	Xaa
7070						7075					7080			
Val	Xaa	Leu	Xaa	Thr	Ser	Gly	Thr	Pro	Xaa	Xaa	Xaa	Pro	Xaa	Xaa
7085						7090					7095			
Thr	Xaa	Xaa	Xaa	Pro	Leu	Leu	Xaa	Pro	Phe	Thr	Leu	Asn	Phe	Thr
7100						7105					7110			
Ile	Thr	Asn	Leu	Xaa	Tyr	Glu	Glu	Xaa	Met	Xaa	Xaa	Pro	Gly	Ser
7115						7120					7125			
Arg	Lys	Phe	Asn	Thr	Thr	Glu	Arg	Val	Leu	Gln	Gly	Leu	Leu	Xaa
7130						7135					7140			
Pro	Xaa	Phe	Lys	Xaa	Thr	Ser	Val	Gly	Xaa	Leu	Tyr	Ser	Gly	Cys
7145						7150					7155			
Arg	Leu	Thr	Leu	Leu	Arg	Xaa	Glu	Lys	Xaa	Xaa	Ala	Ala	Thr	Xaa
7160						7165					7170			
Val	Asp	Xaa	Xaa	Cys	Xaa	Xaa	Xaa	Xaa	Asp	Pro	Xaa	Xaa	Pro	Gly
7175						7180					7185			
Leu	Asp	Arg	Glu	Xaa	Leu	Tyr	Trp	Glu	Leu	Ser	Xaa	Leu	Thr	Asn
7190						7195					7200			
Ser	Ile	Thr	Glu	Leu	Gly	Pro	Tyr	Thr	Leu	Asp	Arg	Asp	Ser	Leu
7205						7210					7215			
Tyr	Val	Asn	Gly	Phe	Thr	His	Arg	Ser	Ser	Met	Pro	Thr	Thr	Ser
7220						7225					7230			
Ile	Pro	Gly	Thr	Ser	Ala	Val	His	Leu	Glu	Thr	Ser	Gly	Thr	Pro
7235						7240					7245			
Ala	Ser	Leu	Pro	Gly	His	Thr	Ala	Pro	Gly	Pro	Leu	Leu	Val	Pro
7250						7255					7260			

Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp
 7265 7270 7275
 Met Arg His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val
 7280 7285 7290
 Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly
 7295 7300 7305
 Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys
 7310 7315 7320
 Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu
 7325 7330 7335
 Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Xaa Leu Tyr Trp Glu
 7340 7345 7350
 Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa
 7355 7360 7365
 Leu Asp Arg Xaa Ser Leu Tyr Val Asn Gly Phe Xaa Xaa Xaa Xaa
 7370 7375 7380
 Xaa Xaa Xaa Xaa Thr Ser Thr Pro Gly Thr Ser Xaa Val Xaa Leu
 7385 7390 7395
 Xaa Thr Ser Gly Thr Pro Xaa Xaa Xaa Pro Xaa Xaa Thr Xaa Xaa
 7400 7405 7410
 Xaa Pro Leu Leu Xaa Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn
 7415 7420 7425
 Leu Xaa Tyr Glu Glu Xaa Met Xaa Xaa Pro Gly Ser Arg Lys Phe
 7430 7435 7440
 Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Xaa Pro Xaa Phe
 7445 7450 7455
 Lys Xaa Thr Ser Val Gly Xaa Leu Tyr Ser Gly Cys Arg Leu Thr
 7460 7465 7470
 Leu Leu Arg Xaa Glu Lys Xaa Xaa Ala Ala Thr Xaa Val Asp Xaa
 7475 7480 7485
 Xaa Cys Xaa Xaa Xaa Xaa Asp Pro Xaa Xaa Pro Gly Leu Asp Arg
 7490 7495 7500
 Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa
 7505 7510 7515
 Glu Leu Gly Pro Tyr Xaa Leu Asp Arg Xaa Ser Leu Tyr Val Asn
 7520 7525 7530
 Gly Phe His Pro Arg Ser Ser Val Pro Thr Thr Ser Thr Pro Gly
 7535 7540 7545
 Thr Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Ser Leu

7550	7555	7560
Pro Gly His Thr Ala	Pro Val Pro Leu Leu Ile	Pro Phe Thr Leu
7565	7570	7575
Asn Phe Thr Ile Thr	Asn Leu His Tyr Glu Glu	Asn Met Gln His
7580	7585	7590
Pro Gly Ser Arg Lys	Phe Asn Thr Thr Glu Arg	Val Leu Gln Gly
7595	7600	7605
Leu Leu Gly Pro Met	Phe Lys Asn Thr Ser	Val Gly Leu Leu Tyr
7610	7615	7620
Ser Gly Cys Arg Leu	Thr Leu Leu Arg Pro	Glu Lys Asn Gly Ala
7625	7630	7635
Ala Thr Gly Met Asp	Ala Ile Cys Ser His	Arg Leu Asp Pro Lys
7640	7645	7650
Ser Pro Gly Leu Asp	Arg Glu Xaa Leu Tyr	Trp Glu Leu Ser Xaa
7655	7660	7665
Leu Thr Xaa Xaa Ile	Xaa Glu Leu Gly Pro	Tyr Xaa Leu Asp Arg
7670	7675	7680
Xaa Ser Leu Tyr Val	Asn Gly Phe Xaa Xaa	Xaa Xaa Xaa Xaa
7685	7690	7695
Xaa Thr Ser Thr Pro	Gly Thr Ser Xaa Val	Xaa Leu Xaa Thr Ser
7700	7705	7710
Gly Thr Pro Xaa Xaa	Xaa Pro Xaa Xaa Thr	Xaa Xaa Xaa Pro Leu
7715	7720	7725
Leu Xaa Pro Phe Thr	Leu Asn Phe Thr Ile	Thr Asn Leu Xaa Tyr
7730	7735	7740
Glu Glu Xaa Met Xaa	Xaa Pro Gly Ser Arg	Lys Phe Asn Thr Thr
7745	7750	7755
Glu Arg Val Leu Gln	Gly Leu Leu Xaa Pro	Xaa Phe Lys Xaa Thr
7760	7765	7770
Ser Val Gly Xaa Leu	Tyr Ser Gly Cys Arg	Leu Thr Leu Leu Arg
7775	7780	7785
Xaa Glu Lys Xaa Xaa	Ala Ala Thr Xaa Val	Asp Xaa Xaa Cys Xaa
7790	7795	7800
Xaa Xaa Xaa Asp Pro	Xaa Xaa Pro Gly Leu	Asp Arg Glu Xaa Leu
7805	7810	7815
Tyr Trp Glu Leu Ser	Xaa Leu Thr Xaa Xaa	Ile Xaa Glu Leu Gly
7820	7825	7830
Pro Tyr Xaa Leu Asp	Arg Xaa Ser Leu Tyr	Val Asn Gly Phe Thr
7835	7840	7845

His Gln Asn Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr	7850	7855	7860
Val Tyr Trp Ala Thr Thr Gly Thr Pro Ser Ser Phe Pro Gly His	7865	7870	7875
Thr Glu Pro Gly Pro Leu Leu Ile Pro Phe Thr Phe Asn Phe Thr	7880	7885	7890
Ile Thr Asn Leu His Tyr Glu Glu Asn Met Gln His Pro Gly Ser	7895	7900	7905
Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Thr	7910	7915	7920
Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys	7925	7930	7935
Arg Leu Thr Leu Leu Arg Pro Glu Lys Gln Glu Ala Ala Thr Gly	7940	7945	7950
Val Asp Thr Ile Cys Thr His Arg Val Asp Pro Ile Gly Pro Gly	7955	7960	7965
Leu Asp Arg Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa	7970	7975	7980
Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa Leu Asp Arg Xaa Ser Leu	7985	7990	7995
Tyr Val Asn Gly Phe Xaa Xaa Xaa Xaa Xaa Xaa Thr Ser	8000	8005	8010
Thr Pro Gly Thr Ser Xaa Val Xaa Leu Xaa Thr Ser Gly Thr Pro	8015	8020	8025
Xaa Xaa Xaa Pro Xaa Xaa Thr Xaa Xaa Xaa Pro Leu Leu Xaa Pro	8030	8035	8040
Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Xaa Tyr Glu Glu Xaa	8045	8050	8055
Met Xaa Xaa Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val	8060	8065	8070
Leu Gln Gly Leu Leu Xaa Pro Xaa Phe Lys Xaa Thr Ser Val Gly	8075	8080	8085
Xaa Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Xaa Glu Lys	8090	8095	8100
Xaa Xaa Ala Ala Thr Xaa Val Asp Xaa Xaa Cys Xaa Xaa Xaa Xaa	8105	8110	8115
Asp Pro Xaa Xaa Pro Gly Leu Asp Arg Glu Xaa Leu Tyr Trp Glu	8120	8125	8130
Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa	8135	8140	8145

Leu Asp Arg Xaa Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser
 8150 8155 8160
 Ser Val Pro Thr Thr Ser Ser Pro Gly Thr Ser Thr Val His Leu
 8165 8170 8175
 Ala Thr Ser Gly Thr Pro Ser Ser Leu Pro Gly His Thr Ala Pro
 8180 8185 8190
 Val Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn
 8195 8200 8205
 Leu His Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe
 8210 8215 8220
 Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe
 8225 8230 8235
 Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr
 8240 8245 8250
 Leu Leu Arg Pro Glu Lys His Gly Ala Ala Thr Gly Val Asp Ala
 8255 8260 8265
 Ile Cys Thr Leu Arg Leu Asp Pro Thr Gly Pro Gly Leu Asp Arg
 8270 8275 8280
 Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa
 8285 8290 8295
 Glu Leu Gly Pro Tyr Xaa Leu Asp Arg Xaa Ser Leu Tyr Val Asn
 8300 8305 8310
 Gly Phe Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Thr Ser Thr Pro Gly
 8315 8320 8325
 Thr Ser Xaa Val Xaa Leu Xaa Thr Ser Gly Thr Pro Xaa Xaa Xaa
 8330 8335 8340
 Pro Xaa Xaa Thr Xaa Xaa Xaa Pro Leu Leu Xaa Pro Phe Thr Leu
 8345 8350 8355
 Asn Phe Thr Ile Thr Asn Leu Xaa Tyr Glu Glu Xaa Met Xaa Xaa
 8360 8365 8370
 Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly
 8375 8380 8385
 Leu Leu Xaa Pro Xaa Phe Lys Xaa Thr Ser Val Gly Xaa Leu Tyr
 8390 8395 8400
 Ser Gly Cys Arg Leu Thr Leu Leu Arg Xaa Glu Lys Xaa Xaa Ala
 8405 8410 8415
 Ala Thr Xaa Val Asp Xaa Xaa Cys Xaa Xaa Xaa Xaa Asp Pro Xaa
 8420 8425 8430
 Xaa Pro Gly Leu Asp Arg Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa

8435		8440		8445
Leu Thr	Xaa Xaa Ile Xaa Glu	Leu Gly Pro Tyr Xaa	Leu Asp Arg	
8450	8455		8460	
Xaa Ser	Leu Tyr Val Asn Gly	Phe Thr His Arg Thr	Ser Val Pro	
8465	8470		8475	
Thr Thr	Ser Thr Pro Gly Thr	Ser Thr Val His Leu	Ala Thr Ser	
8480	8485		8490	
Gly Thr	Pro Ser Ser Leu Pro	Gly His Thr Ala Pro	Val Pro Leu	
8495	8500		8505	
Leu Ile	Pro Phe Thr Leu Asn	Phe Thr Ile Thr Asn	Leu Gln Tyr	
8510	8515		8520	
Glu Glu	Asp Met His Arg Pro	Gly Ser Arg Lys Phe	Asn Thr Thr	
8525	8530		8535	
Glu Arg	Val Leu Gln Gly Leu	Leu Ser Pro Ile Phe	Lys Asn Ser	
8540	8545		8550	
Ser Val	Gly Pro Leu Tyr Ser	Gly Cys Arg Leu Thr	Ser Leu Arg	
8555	8560		8565	
Pro Glu	Lys Asp Gly Ala Ala	Thr Gly Met Asp Ala	Val Cys Leu	
8570	8575		8580	
Tyr His	Pro Asn Pro Lys Arg	Pro Gly Leu Asp Arg	Glu Gln Leu	
8585	8590		8595	
Tyr Cys	Glu Leu Ser Gln Leu	Thr His Asn Ile Thr	Glu Leu Gly	
8600	8605		8610	
Pro Tyr	Ser Leu Asp Arg Asp	Ser Leu Tyr Val Asn	Gly Phe Thr	
8615	8620		8625	
His Gln	Asn Ser Val Pro Thr	Thr Ser Thr Pro Gly	Thr Ser Thr	
8630	8635		8640	
Val Tyr	Trp Ala Thr Thr Gly	Thr Pro Ser Ser Phe	Pro Gly His	
8645	8650		8655	
Thr Xaa	Xaa Xaa Pro Leu Leu	Xaa Pro Phe Thr Leu	Asn Phe Thr	
8660	8665		8670	
Ile Thr	Asn Leu Xaa Tyr Glu	Glu Xaa Met Xaa Xaa	Pro Gly Ser	
8675	8680		8685	
Arg Lys	Phe Asn Thr Thr Glu	Arg Val Leu Gln Gly	Leu Leu Xaa	
8690	8695		8700	
Pro Xaa	Phe Lys Xaa Thr Ser	Val Gly Xaa Leu Tyr	Ser Gly Cys	
8705	8710		8715	
Arg Leu	Thr Leu Leu Arg Xaa	Glu Lys Xaa Xaa Ala	Ala Thr Xaa	
8720	8725		8730	

Val Asp	Xaa Xaa Cys Xaa Xaa	Xaa Xaa Asp Pro Xaa	Xaa Pro Gly
8735		8740	8745
Leu Asp	Arg Glu Xaa Leu Tyr	Trp Glu Leu Ser Xaa	Leu Thr Xaa
8750		8755	8760
Xaa Ile	Xaa Glu Leu Gly Pro	Tyr Xaa Leu Asp Arg	Xaa Ser Leu
8765		8770	8775
Tyr Val	Asn Gly Phe Thr His	Trp Ser Ser Gly Leu	Thr Thr Ser
8780		8785	8790
Thr Pro	Trp Thr Ser Thr Val	Asp Leu Gly Thr Ser	Gly Thr Pro
8795		8800	8805
Ser Pro	Val Pro Ser Pro Thr	Thr Ala Gly Pro Leu	Leu Val Pro
8810		8815	8820
Phe Thr	Leu Asn Phe Thr Ile	Thr Asn Leu Gln Tyr	Glu Glu Asp
8825		8830	8835
Met His	Arg Pro Gly Ser Arg	Lys Phe Asn Ala Thr	Glu Arg Val
8840		8845	8850
Leu Gln	Gly Leu Leu Ser Pro	Ile Phe Lys Asn Thr	Ser Val Gly
8855		8860	8865
Pro Leu	Tyr Ser Gly Cys Arg	Leu Thr Leu Leu Arg	Pro Glu Lys
8870		8875	8880
Gln Glu	Ala Ala Thr Gly Val	Asp Thr Ile Cys Thr	His Arg Val
8885		8890	8895
Asp Pro	Ile Gly Pro Gly Leu	Asp Arg Glu Xaa Leu	Tyr Trp Glu
8900		8905	8910
Leu Ser	Xaa Leu Thr Xaa Xaa	Ile Xaa Glu Leu Gly	Pro Tyr Xaa
8915		8920	8925
Leu Asp	Arg Xaa Ser Leu Tyr	Val Asn Gly Phe Xaa	Xaa Xaa Xaa
8930		8935	8940
Xaa Xaa	Xaa Xaa Thr Ser Thr	Pro Gly Thr Ser Xaa	Val Xaa Leu
8945		8950	8955
Xaa Thr	Ser Gly Thr Pro Xaa	Xaa Xaa Pro Xaa Xaa	Thr Xaa Xaa
8960		8965	8970
Xaa Pro	Leu Leu Xaa Pro Phe	Thr Leu Asn Phe Thr	Ile Thr Asn
8975		8980	8985
Leu Xaa	Tyr Glu Glu Xaa Met	Xaa Xaa Pro Gly Ser	Arg Lys Phe
8990		8995	9000
Asn Thr	Thr Glu Arg Val Leu	Gln Gly Leu Leu Xaa	Pro Xaa Phe
9005		9010	9015
Lys Xaa	Thr Ser Val Gly Xaa	Leu Tyr Ser Gly Cys	Arg Leu Thr
9020		9025	9030

Leu	Leu	Arg	Xaa	Glu	Lys	Xaa	Xaa	Ala	Ala	Thr	Xaa	Val	Asp	Xaa
9035						9040					9045			
Xaa	Cys	Xaa	Xaa	Xaa	Xaa	Asp	Pro	Xaa	Xaa	Pro	Gly	Leu	Asp	Arg
9050						9055					9060			
Glu	Xaa	Leu	Tyr	Trp	Glu	Leu	Ser	Xaa	Leu	Thr	Xaa	Xaa	Ile	Xaa
9065						9070					9075			
Glu	Leu	Gly	Pro	Tyr	Xaa	Leu	Asp	Arg	Xaa	Ser	Leu	Tyr	Val	Asn
9080						9085					9090			
Gly	Phe	Thr	His	Arg	Ser	Phe	Gly	Leu	Thr	Thr	Ser	Thr	Pro	Trp
9095						9100					9105			
Thr	Ser	Thr	Val	Asp	Leu	Gly	Thr	Ser	Gly	Thr	Pro	Ser	Pro	Val
9110						9115					9120			
Pro	Ser	Pro	Thr	Thr	Ala	Gly	Pro	Leu	Leu	Val	Pro	Phe	Thr	Leu
9125						9130					9135			
Asn	Phe	Thr	Ile	Thr	Asn	Leu	Gln	Tyr	Glu	Glu	Asp	Met	His	Arg
9140						9145					9150			
Pro	Gly	Ser	Arg	Lys	Phe	Asn	Thr	Thr	Glu	Arg	Val	Leu	Gln	Gly
9155						9160					9165			
Leu	Leu	Thr	Pro	Leu	Phe	Arg	Asn	Thr	Ser	Val	Ser	Ser	Leu	Tyr
9170						9175					9180			
Ser	Gly	Cys	Arg	Leu	Thr	Leu	Leu	Arg	Pro	Glu	Lys	Asp	Gly	Ala
9185						9190					9195			
Ala	Thr	Arg	Val	Asp	Ala	Val	Cys	Thr	His	Arg	Pro	Asp	Pro	Lys
9200						9205					9210			
Ser	Pro	Gly	Leu	Asp	Arg	Glu	Xaa	Leu	Tyr	Trp	Glu	Leu	Ser	Xaa
9215						9220					9225			
Leu	Thr	Xaa	Xaa	Ile	Xaa	Glu	Leu	Gly	Pro	Tyr	Xaa	Leu	Asp	Arg
9230						9235					9240			
Xaa	Ser	Leu	Tyr	Val	Asn	Gly	Phe	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa
9245						9250					9255			
Xaa	Thr	Ser	Thr	Pro	Gly	Thr	Ser	Xaa	Val	Xaa	Leu	Xaa	Thr	Ser
9260						9265					9270			
Gly	Thr	Pro	Xaa	Xaa	Xaa	Pro	Xaa	Xaa	Thr	Xaa	Xaa	Xaa	Pro	Leu
9275						9280					9285			
Leu	Xaa	Pro	Phe	Thr	Leu	Asn	Phe	Thr	Ile	Thr	Asn	Leu	Xaa	Tyr
9290						9295					9300			
Glu	Glu	Xaa	Met	Xaa	Xaa	Pro	Gly	Ser	Arg	Lys	Phe	Asn	Thr	Thr
9305						9310					9315			
Glu	Arg	Val	Leu	Gln	Gly	Leu	Leu	Xaa	Pro	Xaa	Phe	Lys	Xaa	Thr

9320					9325					9330				
Ser	Val	Gly	Xaa	Leu	Tyr	Ser	Gly	Cys	Arg	Leu	Thr	Leu	Leu	Arg
9335						9340					9345			
Xaa	Glu	Lys	Xaa	Xaa	Ala	Ala	Thr	Xaa	Val	Asp	Xaa	Xaa	Cys	Xaa
9350					9355						9360			
Xaa	Xaa	Xaa	Asp	Pro	Xaa	Xaa	Pro	Gly	Leu	Asp	Arg	Glu	Xaa	Leu
9365					9370						9375			
Tyr	Trp	Glu	Leu	Ser	Xaa	Leu	Thr	Xaa	Xaa	Ile	Xaa	Glu	Leu	Gly
9380					9385						9390			
Pro	Tyr	Xaa	Leu	Asp	Arg	Xaa	Ser	Leu	Tyr	Val	Asn	Gly	Phe	Thr
9395					9400						9405			
His	Trp	Ile	Pro	Val	Pro	Thr	Ser	Ser	Thr	Pro	Gly	Thr	Ser	Thr
9410					9415						9420			
Val	Asp	Leu	Gly	Ser	Gly	Thr	Pro	Ser	Ser	Leu	Pro	Ser	Pro	Thr
9425					9430						9435			
Thr	Ala	Gly	Pro	Leu	Leu	Val	Pro	Phe	Thr	Leu	Asn	Phe	Thr	Ile
9440					9445						9450			
Thr	Asn	Leu	Gln	Tyr	Gly	Glu	Asp	Met	Gly	His	Pro	Gly	Ser	Arg
9455					9460						9465			
Lys	Phe	Asn	Thr	Thr	Glu	Arg	Val	Leu	Gln	Gly	Leu	Leu	Gly	Pro
9470					9475						9480			
Ile	Phe	Lys	Asn	Thr	Ser	Val	Gly	Pro	Leu	Tyr	Ser	Gly	Cys	Arg
9485					9490						9495			
Leu	Thr	Ser	Leu	Arg	Ser	Glu	Lys	Asp	Gly	Ala	Ala	Thr	Gly	Val
9500					9505						9510			
Asp	Ala	Ile	Cys	Ile	His	His	Leu	Asp	Pro	Lys	Ser	Pro	Gly	Leu
9515					9520						9525			
Asp	Arg	Glu	Xaa	Leu	Tyr	Trp	Glu	Leu	Ser	Xaa	Leu	Thr	Xaa	Xaa
9530					9535						9540			
Ile	Xaa	Glu	Leu	Gly	Pro	Tyr	Xaa	Leu	Asp	Arg	Xaa	Ser	Leu	Tyr
9545					9550						9555			
Val	Asn	Gly	Phe	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Thr	Ser	Thr	
9560					9565						9570			
Pro	Gly	Thr	Ser	Xaa	Val	Xaa	Leu	Xaa	Thr	Ser	Gly	Thr	Pro	Xaa
9575					9580						9585			
Xaa	Xaa	Pro	Xaa	Xaa	Thr	Xaa	Xaa	Pro	Leu	Leu	Xaa	Pro	Phe	
9590					9595						9600			
Thr	Leu	Asn	Phe	Thr	Ile	Thr	Asn	Leu	Xaa	Tyr	Glu	Glu	Xaa	Met
9605					9610						9615			

Xaa	Xaa	Pro	Gly	Ser	Arg	Lys	Phe	Asn	Thr	Thr	Glu	Arg	Val	Leu
9620						9625					9630			
Gln	Gly	Leu	Leu	Xaa	Pro	Xaa	Phe	Lys	Xaa	Thr	Ser	Val	Gly	Xaa
9635						9640					9645			
Leu	Tyr	Ser	Gly	Cys	Arg	Leu	Thr	Leu	Leu	Arg	Xaa	Glu	Lys	Xaa
9650						9655					9660			
Xaa	Ala	Ala	Thr	Xaa	Val	Asp	Xaa	Xaa	Cys	Xaa	Xaa	Xaa	Xaa	Asp
9665						9670					9675			
Pro	Xaa	Xaa	Pro	Gly	Leu	Asp	Arg	Glu	Xaa	Leu	Tyr	Trp	Glu	Leu
9680						9685					9690			
Ser	Xaa	Leu	Thr	Xaa	Xaa	Ile	Xaa	Glu	Leu	Gly	Pro	Tyr	Xaa	Leu
9695						9700					9705			
Asp	Arg	Xaa	Ser	Leu	Tyr	Val	Asn	Gly	Phe	Thr	His	Gln	Thr	Phe
9710						9715					9720			
Ala	Pro	Asn	Thr	Ser	Thr	Pro	Gly	Thr	Ser	Thr	Val	Asp	Leu	Gly
9725						9730					9735			
Thr	Ser	Gly	Thr	Pro	Ser	Ser	Leu	Pro	Ser	Pro	Thr	Ser	Ala	Gly
9740						9745					9750			
Pro	Leu	Leu	Val	Pro	Phe	Thr	Leu	Asn	Phe	Thr	Ile	Thr	Asn	Leu
9755						9760					9765			
Gln	Tyr	Glu	Glu	Asp	Met	His	His	Pro	Gly	Ser	Arg	Lys	Phe	Asn
9770						9775					9780			
Thr	Thr	Glu	Arg	Val	Leu	Gln	Gly	Leu	Leu	Gly	Pro	Met	Phe	Lys
9785						9790					9795			
Asn	Thr	Ser	Val	Gly	Leu	Leu	Tyr	Ser	Gly	Cys	Arg	Leu	Thr	Leu
9800						9805					9810			
Leu	Arg	Pro	Glu	Lys	Asn	Gly	Ala	Ala	Thr	Arg	Val	Asp	Ala	Val
9815						9820					9825			
Cys	Thr	His	Arg	Pro	Asp	Pro	Lys	Ser	Pro	Gly	Leu	Asp	Arg	Glu
9830						9835					9840			
Xaa	Leu	Tyr	Trp	Glu	Leu	Ser	Xaa	Leu	Thr	Xaa	Xaa	Ile	Xaa	Glu
9845						9850					9855			
Leu	Gly	Pro	Tyr	Xaa	Leu	Asp	Arg	Xaa	Ser	Leu	Tyr	Val	Asn	Gly
9860						9865					9870			
Phe	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Thr	Ser	Thr	Pro	Gly	Thr
9875						9880					9885			
Ser	Xaa	Val	Xaa	Leu	Xaa	Thr	Ser	Gly	Thr	Pro	Xaa	Xaa	Xaa	Pro
9890						9895					9900			
Xaa	Xaa	Thr	Ala	Pro	Val	Pro	Leu	Leu	Ile	Pro	Phe	Thr	Leu	Asn
9905						9910					9915			

Phe Thr	Ile Thr Asn Leu His	Tyr Glu Glu Asn Met	Gln His Pro
9920	9925	9930	
Gly Ser	Arg Lys Phe Asn Thr	Thr Glu Arg Val Leu	Gln Gly Leu
9935	9940	9945	
Leu Arg	Pro Leu Phe Lys Ser	Thr Ser Val Gly Pro	Leu Tyr Ser
9950	9955	9960	
Gly Cys	Arg Leu Thr Leu Leu	Arg Pro Glu Lys His	Gly Ala Ala
9965	9970	9975	
Thr Gly	Val Asp Ala Ile Cys	Thr Leu Arg Leu Asp	Pro Thr Gly
9980	9985	9990	
Pro Gly	Leu Asp Arg Glu Arg	Leu Tyr Trp Glu Leu	Ser Gln Leu
9995	10000	10005	
Thr Asn	Ser Val Thr Glu Leu	Gly Pro Tyr Thr Leu	Asp Arg Asp
10010	10015	10020	
Ser Leu	Tyr Val Asn Gly Phe	Thr Gln Arg Ser Ser	Val Pro Thr
10025	10030	10035	
Thr Ser	Ile Pro Gly Thr Ser	Ala Val His Leu Glu	Thr Ser Gly
10040	10045	10050	
Thr Pro	Ala Ser Leu Pro Gly	His Thr Ala Pro Gly	Pro Leu Leu
10055	10060	10065	
Val Pro	Phe Thr Leu Asn Phe	Thr Ile Thr Asn Leu	Gln Tyr Glu
10070	10075	10080	
Val Asp	Met Arg His Pro Gly	Ser Arg Lys Phe Asn	Thr Thr Glu
10085	10090	10095	
Arg Val	Leu Gln Gly Leu Leu	Lys Pro Leu Phe Lys	Ser Thr Ser
10100	10105	10110	
Val Gly	Pro Leu Tyr Ser Gly	Cys Arg Leu Thr Leu	Leu Arg Pro
10115	10120	10125	
Glu Lys	Arg Gly Ala Ala Thr	Gly Val Asp Thr Ile	Cys Thr His
10130	10135	10140	
Arg Leu	Asp Pro Leu Asn Pro	Gly Leu Asp Arg Glu	Gln Leu Tyr
10145	10150	10155	
Trp Glu	Leu Ser Lys Leu Thr	Arg Gly Ile Ile Glu	Leu Gly Pro
10160	10165	10170	
Tyr Leu	Leu Asp Arg Gly Ser	Leu Tyr Val Asn Gly	Phe Thr His
10175	10180	10185	
Arg Asn	Phe Val Pro Ile Thr	Ser Thr Pro Gly Thr	Ser Thr Val
10190	10195	10200	
His Leu	Gly Thr Ser Glu Thr	Pro Ser Ser Leu Pro	Arg Pro Ile

10205	10210	10215
Val Pro Gly Pro Leu Leu	Val Pro Phe Thr Leu	Asn Phe Thr Ile
10220	10225	10230
Thr Asn Leu Gln Tyr Glu	Glu Ala Met Arg His	Pro Gly Ser Arg
10235	10240	10245
Lys Phe Asn Thr Thr Glu	Arg Val Leu Gln Gly	Leu Leu Arg Pro
10250	10255	10260
Leu Phe Lys Asn Thr Ser	Ile Gly Pro Leu Tyr	Ser Ser Cys Arg
10265	10270	10275
Leu Thr Leu Leu Arg Pro	Glu Lys Asp Lys Ala	Ala Thr Arg Val
10280	10285	10290
Asp Ala Ile Cys Thr His	His Pro Asp Pro Gln	Ser Pro Gly Leu
10295	10300	10305
Asn Arg Glu Gln Leu Tyr	Trp Glu Leu Ser Gln	Leu Thr His Gly
10310	10315	10320
Ile Thr Glu Leu Gly Pro	Tyr Thr Leu Asp Arg	Asp Ser Leu Tyr
10325	10330	10335
Val Asp Gly Phe Thr His	Trp Ser Pro Ile Pro	Thr Thr Ser Thr
10340	10345	10350
Pro Gly Thr Ser Ile Val	Asn Leu Gly Thr Ser	Gly Ile Pro Pro
10355	10360	10365
Ser Leu Pro Glu Thr Thr	Xaa Xaa Xaa Pro Leu	Leu Xaa Pro Phe
10370	10375	10380
Thr Leu Asn Phe Thr Ile	Thr Asn Leu Xaa Tyr	Glu Glu Xaa Met
10385	10390	10395
Xaa Xaa Pro Gly Ser Arg	Lys Phe Asn Thr Thr	Glu Arg Val Leu
10400	10405	10410
Gln Gly Leu Leu Lys Pro	Leu Phe Lys Ser Thr	Ser Val Gly Pro
10415	10420	10425
Leu Tyr Ser Gly Cys Arg	Leu Thr Leu Leu Arg	Pro Glu Lys Asp
10430	10435	10440
Gly Val Ala Thr Arg Val	Asp Ala Ile Cys Thr	His Arg Pro Asp
10445	10450	10455
Pro Lys Ile Pro Gly Leu	Asp Arg Gln Gln Leu	Tyr Trp Glu Leu
10460	10465	10470
Ser Gln Leu Thr His Ser	Ile Thr Glu Leu Gly	Pro Tyr Thr Leu
10475	10480	10485
Asp Arg Asp Ser Leu Tyr	Val Asn Gly Phe Thr	Gln Arg Ser Ser
10490	10495	10500

Val Pro	Thr Thr	Ser Thr	Pro	Gly Thr	Phe Thr	Val	Gln Pro	Glu
10505			10510			10515		
Thr Ser	Glu Thr	Pro Ser	Ser	Leu Pro	Gly Pro	Thr	Ala Thr	Gly
10520			10525			10530		
Pro Val	Leu Leu	Pro Phe	Thr	Leu Asn	Phe Thr	Ile	Thr Asn	Leu
10535			10540			10545		
Gln Tyr	Glu Glu	Asp Met	His	Arg Pro	Gly Ser	Arg	Lys Phe	Asn
10550			10555			10560		
Thr Thr	Glu Arg	Val Leu	Gln	Gly Leu	Leu Met	Pro	Leu Phe	Lys
10565			10570			10575		
Asn Thr	Ser Val	Ser Ser	Leu	Tyr Ser	Gly Cys	Arg	Leu Thr	Leu
10580			10585			10590		
Leu Arg	Pro Glu	Lys Asp	Gly	Ala Ala	Thr Arg	Val	Asp Ala	Val
10595			10600			10605		
Cys Thr	His Arg	Pro Asp	Pro	Lys Ser	Pro Gly	Leu	Asp Arg	Glu
10610			10615			10620		
Arg Leu	Tyr Trp	Lys Leu	Ser	Gln Leu	Thr His	Gly	Ile Thr	Glu
10625			10630			10635		
Leu Gly	Pro Tyr	Thr Leu	Asp	Arg His	Ser Leu	Tyr	Val Asn	Gly
10640			10645			10650		
Phe Thr	His Gln	Ser Ser	Met	Thr Thr	Thr Arg	Thr	Pro Asp	Thr
10655			10660			10665		
Ser Thr	Met His	Leu Ala	Thr	Ser Arg	Thr Pro	Ala	Ser Leu	Ser
10670			10675			10680		
Gly Pro	Thr Thr	Ala Ser	Pro	Leu Leu	Val Leu	Phe	Thr Ile	Asn
10685			10690			10695		
Phe Thr	Ile Thr	Asn Leu	Arg	Tyr Glu	Glu Asn	Met	His His	Pro
10700			10705			10710		
Gly Ser	Arg Lys	Phe Asn	Thr	Thr Glu	Arg Val	Leu	Gln Gly	Leu
10715			10720			10725		
Leu Arg	Pro Val	Phe Lys	Asn	Thr Ser	Val Gly	Pro	Leu Tyr	Ser
10730			10735			10740		
Gly Cys	Arg Leu	Thr Leu	Leu	Arg Pro	Lys Lys	Asp	Gly Ala	Ala
10745			10750			10755		
Thr Lys	Val Asp	Ala Ile	Cys	Thr Tyr	Arg Pro	Asp	Pro Lys	Ser
10760			10765			10770		
Pro Gly	Leu Asp	Arg Glu	Gln	Leu Tyr	Trp Glu	Leu	Ser Gln	Leu
10775			10780			10785		
Thr His	Ser Ile	Thr Glu	Leu	Gly Pro	Tyr Thr	Gln	Asp Arg	Asp
10790			10795			10800		

Ser Leu	Tyr Asn Val Gly	Phe	Thr Gln Arg Ser Ser	Val Pro Thr
10805		10810		10815
Thr Ser	Val Pro Gly Thr	Pro	Thr Val Asp Leu Gly	Thr Ser Gly
10820		10825		10830
Thr Pro	Val Ser Lys Pro Gly	Pro Ser Ala Ala Ser	Pro Leu Leu	
10835		10840		10845
Val Leu	Phe Thr Leu Asn Gly	Thr Ile Thr Asn Leu	Arg Tyr Glu	
10850		10855		10860
Glu Asn	Met Gln His Pro Gly	Ser Arg Lys Phe Asn	Thr Thr Glu	
10865		10870		10875
Arg Val	Leu Gln Gly Leu Leu	Arg Ser Leu Phe Lys	Ser Thr Ser	
10880		10885		10890
Val Gly	Pro Leu Tyr Ser Gly	Cys Arg Leu Thr Leu	Leu Arg Pro	
10895		10900		10905
Glu Lys	Asp Gly Thr Ala Thr	Gly Val Asp Ala Ile	Cys Thr His	
10910		10915		10920
His Pro	Asp Pro Lys Ser Pro	Arg Leu Asp Arg Glu	Gln Leu Tyr	
10925		10930		10935
Trp Glu	Leu Ser Gln Leu Thr	His Asn Ile Thr Glu	Leu Gly His	
10940		10945		10950
Tyr Ala	Leu Asp Asn Asp Ser	Leu Phe Val Asn Gly	Phe Thr His	
10955		10960		10965
Arg Ser	Ser Val Ser Thr Thr	Ser Thr Pro Gly Thr	Pro Thr Val	
10970		10975		10980
Tyr Leu	Gly Ala Ser Lys Thr	Pro Ala Ser Ile Phe	Gly Pro Ser	
10985		10990		10995
Ala Ala	Ser His Leu Leu Ile	Leu Phe Thr Leu Asn	Phe Thr Ile	
11000		11005		11010
Thr Asn	Leu Arg Tyr Glu Glu	Asn Met Trp Pro Gly	Ser Arg Lys	
11015		11020		11025
Phe Asn	Thr Thr Glu Arg Val	Leu Gln Gly Leu Leu	Arg Pro Leu	
11030		11035		11040
Phe Lys	Asn Thr Ser Val Gly	Pro Leu Tyr Ser Gly	Ser Arg Leu	
11045		11050		11055
Thr Leu	Leu Arg Pro Glu Lys	Asp Gly Glu Ala Thr	Gly Val Asp	
11060		11065		11070
Ala Ile	Cys Thr His Arg Pro	Asp Pro Thr Gly Pro	Gly Leu Asp	
11075		11080		11085
Arg Glu	Gln Leu Tyr Leu Glu	Leu Ser Gln Leu Thr	His Ser Ile	

11090					11095					11100				
Thr	Glu	Leu	Gly	Pro	Tyr	Thr	Leu	Asp	Arg	Asp	Ser	Leu	Tyr	Val
11105						11110					11115			
Asn	Gly	Phe	Thr	His	Arg	Ser	Ser	Val	Pro	Thr	Thr	Ser	Thr	Gly
11120						11125					11130			
Val	Val	Ser	Glu	Glu	Pro	Phe	Thr	Leu	Asn	Phe	Thr	Ile	Asn	Asn
11135						11140					11145			
Leu	Arg	Tyr	Met	Ala	Asp	Met	Gly	Gln	Pro	Gly	Ser	Leu	Lys	Phe
11150						11155					11160			
Asn	Ile	Thr	Asp	Asn	Val	Met	Lys	His	Leu	Leu	Ser	Pro	Leu	Phe
11165						11170					11175			
Gln	Arg	Ser	Ser	Leu	Gly	Ala	Arg	Tyr	Thr	Gly	Cys	Arg	Val	Ile
11180						11185					11190			
Ala	Leu	Arg	Ser	Val	Lys	Asn	Gly	Ala	Glu	Thr	Arg	Val	Asp	Leu
11195						11200					11205			
Leu	Cys	Thr	Tyr	Leu	Gln	Pro	Leu	Ser	Gly	Pro	Gly	Leu	Pro	Ile
11210						11215					11220			
Lys	Gln	Val	Phe	His	Glu	Leu	Ser	Gln	Gln	Thr	His	Gly	Ile	Thr
11225						11230					11235			
Arg	Leu	Gly	Pro	Tyr	Ser	Leu	Asp	Lys	Asp	Ser	Leu	Tyr	Leu	Asn
11240						11245					11250			
Gly	Tyr	Asn	Glu	Pro	Gly	Leu	Asp	Glu	Pro	Pro	Thr	Thr	Pro	Lys
11255						11260					11265			
Pro	Ala	Thr	Thr	Phe	Leu	Pro	Pro	Leu	Ser	Glu	Ala	Thr	Thr	Ala
11270						11275					11280			
Met	Gly	Tyr	His	Leu	Lys	Thr	Leu	Thr	Leu	Asn	Phe	Thr	Ile	Ser
11285						11290					11295			
Asn	Leu	Gln	Tyr	Ser	Pro	Asp	Met	Gly	Lys	Gly	Ser	Ala	Thr	Phe
11300						11305					11310			
Asn	Ser	Thr	Glu	Gly	Val	Leu	Gln	His	Leu	Leu	Arg	Pro	Leu	Phe
11315						11320					11325			
Gln	Lys	Ser	Ser	Met	Gly	Pro	Phe	Tyr	Leu	Gly	Cys	Gln	Leu	Ile
11330						11335					11340			
Ser	Leu	Arg	Pro	Glu	Lys	Asp	Gly	Ala	Ala	Thr	Gly	Val	Asp	Thr
11345						11350					11355			
Thr	Cys	Thr	Tyr	His	Pro	Asp	Pro	Val	Gly	Pro	Gly	Leu	Asp	Ile
11360						11365					11370			
Gln	Gln	Leu	Tyr	Trp	Glu	Leu	Ser	Gln	Leu	Thr	His	Gly	Val	Thr
11375						11380					11385			

Gln	Leu	Gly	Phe	Tyr	Val	Leu	Asp	Arg	Asp	Ser	Leu	Phe	Ile	Asn
11390						11395					11400			
Gly	Tyr	Ala	Pro	Gln	Asn	Leu	Ser	Ile	Arg	Gly	Glu	Tyr	Gln	Ile
11405						11410					11415			
Asn	Phe	His	Ile	Val	Asn	Trp	Asn	Leu	Ser	Asn	Pro	Asp	Pro	Thr
11420						11425					11430			
Ser	Ser	Glu	Tyr	Ile	Thr	Leu	Leu	Arg	Asp	Ile	Gln	Asp	Lys	Val
11435						11440					11445			
Thr	Thr	Leu	Tyr	Lys	Gly	Ser	Gln	Leu	His	Asp	Thr	Phe	Arg	Phe
11450						11455					11460			
Cys	Leu	Val	Thr	Asn	Leu	Thr	Met	Asp	Ser	Val	Leu	Val	Thr	Val
11465						11470					11475			
Lys	Ala	Leu	Phe	Ser	Ser	Asn	Leu	Asp	Pro	Ser	Leu	Val	Glu	Gln
11480						11485					11490			
Val	Phe	Leu	Asp	Lys	Thr	Leu	Asn	Ala	Ser	Phe	His	Trp	Leu	Gly
11495						11500					11505			
Ser	Thr	Tyr	Gln	Leu	Val	Asp	Ile	His	Val	Thr	Glu	Met	Glu	Ser
11510						11515					11520			
Ser	Val	Tyr	Gln	Pro	Thr	Ser	Ser	Ser	Ser	Thr	Gln	His	Phe	Tyr
11525						11530					11535			
Leu	Asn	Phe	Thr	Ile	Thr	Asn	Leu	Pro	Tyr	Ser	Gln	Asp	Lys	Ala
11540						11545					11550			
Gln	Pro	Gly	Thr	Thr	Asn	Tyr	Gln	Arg	Asn	Lys	Arg	Asn	Ile	Glu
11555						11560					11565			
Asp	Ala	Leu	Asn	Gln	Leu	Phe	Arg	Asn	Ser	Ser	Ile	Lys	Ser	Tyr
11570						11575					11580			
Phe	Ser	Asp	Cys	Gln	Val	Ser	Thr	Phe	Arg	Ser	Val	Pro	Asn	Arg
11585						11590					11595			
His	His	Thr	Gly	Val	Asp	Ser	Leu	Cys	Asn	Phe	Ser	Pro	Leu	Ala
11600						11605					11610			
Arg	Arg	Val	Asp	Arg	Val	Ala	Ile	Tyr	Glu	Glu	Phe	Leu	Arg	Met
11615						11620					11625			
Thr	Arg	Asn	Gly	Thr	Gln	Leu	Gln	Asn	Phe	Thr	Leu	Asp	Arg	Ser
11630						11635					11640			
Ser	Val	Leu	Val	Asp	Gly	Tyr	Ser	Pro	Asn	Arg	Asn	Glu	Pro	Leu
11645						11650					11655			
Thr	Gly	Asn	Ser	Asp	Leu	Pro	Phe	Trp	Ala	Val	Ile	Leu	Ile	Gly
11660						11665					11670			
Leu	Ala	Gly	Leu	Leu	Gly	Leu	Ile	Thr	Cys	Leu	Ile	Cys	Gly	Val
11675						11680					11685			

Leu Val Thr Thr Arg Arg Arg Lys Lys Glu Gly Glu Tyr Asn Val
11690 11695 11700

Gln Gln Gln Cys Pro Gly Tyr Tyr Gln Ser His Leu Asp Leu Glu
11705 11710 11715

Asp Leu Gln
11720

<210> 163

<211> 156

<212> PRT

<213> Homo sapiens

<400> 163

Thr Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr
1 5 10 15

Asn Leu Gln Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe
20 25 30

Asn Ala Thr Glu Arg Val Leu Gln Gly Leu Leu Ser Pro Ile Phe Lys
35 40 45

Asn Ser Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu
50 55 60

Arg Pro Glu Lys Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu
65 70 75 80

Tyr His Pro Asn Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu Tyr
85 90 95

Trp Glu Leu Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly Pro Tyr
100 105 110

Ser Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Gln Asn
115 120 125

Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Tyr Trp Ala
130 135 140

Thr Thr Gly Thr Pro Ser Ser Phe Pro Gly His Thr
145 150 155

<210> 164

<211> 42

<212> PRT

<213> Homo sapiens

<400> 164

```
Ala Thr Val Pro Phe Met Val Pro Phe Thr Leu Asn Phe Thr Ile Thr
1           5           10           15

Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe
          20           25           30

Asn Ala Thr Thr Glu Arg Glu Leu Gln Gly Leu
          35           40
```

<210> 165

<211> 42

<212> PRT

<213> Homo sapiens

<400> 165

```
Thr Ala Val Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr
1           5           10           15

Asn Leu Gln Tyr Gly Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe
          20           25           30

Asn Thr Thr Thr Glu Arg Val Leu Gln Gly Leu
          35           40
```

<210> 166

<211> 42

<212> PRT

<213> Homo sapiens

<400> 166

```
Val Pro Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr
1           5           10           15

Asn Leu Gln Tyr Glu Glu Ala Met Arg His Pro Gly Ser Arg Lys Phe
          20           25           30

Asn Thr Thr Thr Glu Arg Val Leu Gln Gly Leu
          35           40
```

<210> 167

<211> 42

<212> PRT

<213> Homo sapiens

<400> 167

Ala Pro Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr
1 5 10 15

Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe
20 25 30

Ser Thr Thr Glu Arg Val Leu Gln Gly Leu
35 40

<210> 168

<211> 42

<212> PRT

<213> Homo sapiens

<400> 168

Ala Pro Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr
1 5 10 15

Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe
20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu
35 40

<210> 169

<211> 42

<212> PRT

<213> Homo sapiens

<400> 169

Ala Pro Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr
1 5 10 15

Asn Leu Gln Tyr Glu Val Asp Met Arg His Pro Gly Ser Arg Lys Phe
20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu

```

35                                40

<210> 170

<211> 42

<212> PRT

<213> Homo sapiens

<400> 170

Ser Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr
1                    5                                10          15

Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe
20                    25                    30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu
35                                40

<210> 171

<211> 42

<212> PRT

<213> Homo sapiens

<400> 171

Ala Ala Gly Pro Leu Leu Met Pro Phe Thr Leu Asn Phe Thr Ile Thr
1                    5                                10          15

Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr Gly Ser Arg Lys Phe
20                    25                    30

Asn Thr Met Glu Ser Val Leu Gln Gly Leu
35                                40

<210> 172

<211> 42

<212> PRT

<213> Homo sapiens

<400> 172

Thr Ala Ser Pro Leu Leu Val Leu Phe Thr Ile Asn Cys Thr Ile Thr
1                    5                                10          15

```

Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr Gly Ser Arg Lys Phe
 20 25 30

Asn Thr Met Glu Ser Val Leu Gln Gly Leu
 35 40

<210> 173

<211> 42

<212> PRT

<213> Homo sapiens

<400> 173

Ala Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr
 1 5 10 15

Asn Leu Gln Tyr Gly Glu Asp Met Gly His Pro Gly Ser Arg Lys Phe
 20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu
 35 40

<210> 174

<211> 42

<212> PRT

<213> Homo sapiens

<400> 174

Thr Ala Gly Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr
 1 5 10 15

Asn Leu Gln Tyr Gly Glu Asp Met Gly His Pro Gly Ser Arg Lys Phe
 20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu
 35 40

<210> 175

<211> 42

<212> PRT

<213> Homo sapiens

<400> 175

Thr Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr
1 5 10 15

Asn Leu Gln Tyr Gly Glu Asp Met Gly His Pro Gly Ser Arg Lys Phe
20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu
35 40

<210> 176

<211> 42

<212> PRT

<213> Homo sapiens

<400> 176

Thr Ala Gly Pro Leu Leu Val Leu Phe Thr Leu Asn Phe Thr Ile Thr
1 5 10 15

Asn Leu Lys Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe
20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Thr Leu
35 40

<210> 177

<211> 42

<212> PRT

<213> Homo sapiens

<400> 177

Thr Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr
1 5 10 15

Asn Leu Gln Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe
20 25 30

Asn Ala Thr Glu Arg Val Leu Gln Gly Leu
35 40

<210> 178

<211> 42

<212> PRT

<213> Homo sapiens

<400> 178

```

Thr Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr
1           5              10              15

Asn Leu Gln Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Arg Phe
20          25          30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu
35          40

```

<210> 179

<211> 42

<212> PRT

<213> Homo sapiens

<400> 179

```

Thr Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr
1           5              10              15

Asn Leu Gln Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe
20          25          30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu
35          40

```

<210> 180

<211> 42

<212> PRT

<213> Homo sapiens

<400> 180

```

Ala Pro Val Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr
1           5              10              15

Asn Leu Gln Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe
20          25          30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu
35          40

```

<210> 181

<213> Homo sapiens

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu
35 40

<213> Homo sapiens

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu
35 40

<213> Homo sapiens

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu

```

35                                40

<210> 184

<211> 42

<212> PRT

<213> Homo sapiens

<400> 184

Thr Ala Ser Pro Leu Leu Val Leu Phe Thr Ile Asn Phe Thr Ile Thr
1          5                                10          15

Asn Gln Arg Tyr Glu Glu Asn Met His His Pro Gly Ser Arg Lys Phe
20                                25          30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu
35                                40

<210> 185

<211> 42

<212> PRT

<213> Homo sapiens

<400> 185

Thr Ala Ser Pro Leu Leu Val Leu Phe Thr Ile Asn Phe Thr Ile Thr
1          5                                10          15

Asn Leu Arg Tyr Glu Glu Asn Met His His Pro Gly Ser Arg Lys Phe
20                                25          30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu
35                                40

<210> 186

<211> 42

<212> PRT

<213> Homo sapiens

<400> 186

Glu Pro Gly Pro Leu Leu Ile Pro Phe Thr Phe Asn Phe Thr Ile Thr
1          5                                10          15

```


Asn Thr Thr Glu Arg Val Leu Gln Gly Leu
35 40

<211> 42

<212> PRT

<213> Homo sapiens

<400> 187

Glu Pro Gly Pro Leu Leu Ile Pro Phe Thr Phe Asn Phe Thr Ile Thr
1 5 10 15

Asn Leu Arg Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe
20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu
35 40

<210> 188

<211> 42

<212> PRT

<213> Homo sapiens

<400> 188

Ala Pro Val Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr
1 5 10 15

Asn Leu His Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe
20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu
35 40

<210> 189

<211> 42

<212> PRT

<213> Homo sapiens

<400> 189

<212> PRT

<213> Homo sapiens

<400> 192

```
Ala Ala Ser His Leu Leu Ile Leu Phe Thr Leu Asn Phe Thr Ile Thr
1          5          10          15

Asn Leu Arg Tyr Glu Glu Asn Met Trp Pro Gly Ser Arg Lys Phe Asn
20          25          30

Thr Thr Glu Arg Val Leu Gln Gly Leu
35          40
```

<210> 193

<211> 42

<212> PRT

<213> Homo sapiens

<400> 193

```
Thr Gly Val Val Ser Glu Glu Pro Phe Thr Leu Asn Phe Thr Ile Asn
1          5          10          15

Asn Leu Arg Tyr Met Ala Asp Met Gly Gln Pro Gly Ser Leu Lys Phe
20          25          30

Asn Ile Thr Asp Asn Val Met Lys His Leu
35          40
```

<210> 194

<211> 42

<212> PRT

<213> Homo sapiens

<400> 194

```
Ala Met Gly Tyr His Leu Lys Thr Leu Thr Leu Asn Phe Thr Ile Ser
1          5          10          15

Asn Leu Gln Tyr Ser Pro Asp Met Gly Lys Gly Ser Ala Thr Phe Asn
20          25          30

Ser Thr Glu Gly Val Leu Gln His Leu Leu
35          40
```

<210> 195

<211> 23

<212> PRT

<213> Homo sapiens

<400> 195

Leu Lys Pro Leu Phe Arg Asn Ser Ser Leu Glu Tyr Leu Tyr Ser Gly
1 5 10 15

Cys Arg Leu Ala Ser Leu Arg
20

<210> 196

<211> 23

<212> PRT

<213> Homo sapiens

<400> 196

Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu Tyr Ser Gly
1 5 10 15

Cys Arg Leu Thr Leu Leu Arg
20

<210> 197

<211> 23

<212> PRT

<213> Homo sapiens

<400> 197

Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly
1 5 10 15

Cys Arg Leu Thr Leu Leu Arg
20

<210> 198

<211> 23

<212> PRT

<213> Homo sapiens

<400> 198

Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly
1 5 10 15

Cys Arg Leu Thr Leu Leu Arg
20

<210> 199

<211> 23

<212> PRT

<213> Homo sapiens

<400> 199

Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Ser
1 5 10 15

Cys Arg Leu Thr Leu Leu Arg
20

<210> 200

<211> 23

<212> PRT

<213> Homo sapiens

<400> 200

Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly
1 5 10 15

Cys Arg Leu Thr Ser Leu Arg
20

<210> 201

<211> 23

<212> PRT

<213> Homo sapiens

<400> 201

Cys Arg Leu Thr Ser Leu Arg

20

<210> 205

<211> 23

<212> PRT

<213> Homo sapiens

<400> 205

Leu	Gly	Pro	Leu	Phe	Lys	Asn	Ser	Ser	Val	Gly	Pro	Leu	Tyr	Ser	Gly
1				5					10				15		

Cys	Arg	Leu	Ile	Ser	Leu	Arg
			20			

<210> 206

<211> 23

<212> PRT

<213> Homo sapiens

<400> 206

Leu	Gly	Pro	Leu	Phe	Lys	Asn	Ser	Ser	Val	Asp	Pro	Leu	Tyr	Ser	Gly
1				5					10				15		

Cys	Arg	Leu	Thr	Ser	Leu	Arg
			20			

<210> 207

<211> 23

<212> PRT

<213> Homo sapiens

<400> 207

Leu	Ser	Pro	Ile	Phe	Lys	Asn	Ser	Ser	Val	Gly	Pro	Leu	Tyr	Ser	Gly
1				5					10				15		

Cys	Arg	Leu	Thr	Ser	Leu	Arg
			20			

<210> 208

<211> 23

<212> PRT

<213> Homo sapiens

<400> 208

Leu Ser Pro Ile Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly
1 5 10 15

Cys Arg Leu Thr Leu Leu Arg
20

<210> 209

<211> 23

<212> PRT

<213> Homo sapiens

<400> 209

Leu Ser Pro Leu Phe Gln Arg Ser Ser Leu Gly Ala Arg Tyr Thr Gly
1 5 10 15

Cys Arg Val Ile Ala Leu Arg
20

<210> 210

<211> 23

<212> PRT

<213> Homo sapiens

<400> 210

Leu Arg Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu Tyr Ser Gly
1 5 10 15

Cys Arg Leu Thr Leu Leu Arg
20

<210> 211

<211> 23

<212> PRT

<213> Homo sapiens

<400> 211

Leu Arg Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly
1 5 10 15

Ser Arg Leu Thr Leu Leu Arg
20

<210> 212

<211> 23

<212> PRT

<213> Homo sapiens

<400> 212

Leu Arg Pro Leu Phe Lys Asn Thr Ser Ile Gly Pro Leu Tyr Ser Ser
1 5 10 15

Cys Arg Leu Thr Leu Leu Arg
20

<210> 213

<211> 23

<212> PRT

<213> Homo sapiens

<400> 213

Leu Arg Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly
1 5 10 15

Cys Arg Leu Thr Leu Leu Arg
20

<210> 214

<211> 23

<212> PRT

<213> Homo sapiens

<400> 214

Leu Arg Pro Val Phe Lys Asn Thr Ser Val Gly Leu Leu Tyr Ser Gly
 1 5 10 15

Cys Arg Leu Thr Leu Leu Arg
 20

<210> 215

<211> 23

<212> PRT

<213> Homo sapiens

<400> 215

Leu Arg Pro Val Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly
 1 5 10 15

Cys Arg Leu Thr Leu Leu Arg
 20

<210> 216

<211> 23

<212> PRT

<213> Homo sapiens

<400> 216

Leu Arg Ser Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly
 1 5 10 15

Cys Arg Leu Thr Leu Leu Arg
 20

<210> 217

<211> 23

<212> PRT

<213> Homo sapiens

<400> 217

Leu Arg Ser Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly
 1 5 10 15

Cys Arg Leu Thr Ser Leu Arg

112200 8629600

20

<210> 218

<211> 23

<212> PRT

<213> Homo sapiens

<400> 218

Leu	Thr	Pro	Leu	Phe	Lys	Asn	Thr	Ser	Val	Gly	Pro	Leu	Tyr	Ser	Gly
1				5					10					15	

Cys	Arg	Leu	Thr	Leu	Leu	Arg
			20			

<210> 219

<211> 23

<212> PRT

<213> Homo sapiens

<400> 219

Leu	Thr	Pro	Leu	Phe	Arg	Asn	Thr	Ser	Val	Ser	Ser	Leu	Tyr	Ser	Gly
1				5					10					15	

Cys	Arg	Leu	Thr	Leu	Leu	Arg
			20			

<210> 220

<211> 23

<212> PRT

<213> Homo sapiens

<400> 220

Leu	Met	Pro	Leu	Phe	Lys	Asn	Thr	Ser	Val	Ser	Ser	Leu	Tyr	Ser	Gly
1				5					10					15	

Cys	Arg	Leu	Thr	Leu	Leu	Arg
			20			

<210> 221

<211> 22

<212> PRT

<213> Homo sapiens

<400> 221

Arg Pro Leu Phe Gln Lys Ser Ser Met Gly Pro Phe Tyr Leu Gly Cys
1 5 10 15

Gln Leu Ile Ser Leu Arg
20

<210> 222

<211> 58

<212> PRT

<213> Homo sapiens

<400> 222

Pro Glu Lys Asp Ser Ser Ala Met Ala Val Asp Ala Ile Cys Thr His
1 5 10 15

Arg Pro Asp Pro Glu Asp Leu Gly Leu Asp Arg Glu Arg Leu Tyr Trp
20 25 30

Glu Leu Ser Asn Leu Thr Asn Gly Ile Gln Glu Leu Gly Pro Tyr Thr
35 40 45

Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly
50 55

<210> 223

<211> 58

<212> PRT

<213> Homo sapiens

<400> 223

Pro Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His
1 5 10 15

Arg Leu Asp Pro Lys Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp
20 25 30

Glu Leu Ser Lys Leu Thr Asn Asp Ile Glu Glu Leu Gly Pro Tyr Thr

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35              40              45
Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly
50              55

<210> 224

<211> 58

<212> PRT

<213> Homo sapiens

<400> 224

Pro Lys Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His
1              5              10              15
Arg Leu Asp Pro Lys Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp
20              25
Glu Leu Ser Lys Leu Thr Asn Asp Ile Glu Glu Leu Gly Pro Tyr Thr
35              40              45
Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly
50              55

<210> 225

<211> 58

<212> PRT

<213> Homo sapiens

<400> 225

Pro Glu Lys Asp Gly Thr Ala Thr Gly Val Asp Ala Ile Cys Thr His
1              5              10              15
His Pro Asp Pro Lys Ser Pro Arg Leu Asp Arg Glu Gln Leu Tyr Trp
20              25              30
Glu Leu Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly His Tyr Ala
35              40              45
Leu Asp Asn Asp Ser Leu Phe Val Asn Gly
50              55

<210> 226

<211> 58

<212> PRT

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<213> Homo sapiens

<400> 226

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Pro Glu Lys Asp Gly Glu Ala Thr Gly Val Asp Ala Ile Cys Thr His
1           5           10           15
Arg Pro Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu Gln Leu Tyr Leu
20           25           30
Glu Leu Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr
35           40           45
Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly
50           55

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<210> 227

<211> 58

<212> PRT

<213> Homo sapiens

<400> 227

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Pro Glu Lys Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu Tyr
1           5           10           15
His Pro Asn Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp
20           25           30
Glu Leu Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly Pro Tyr Ser
35           40           45
Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly
50           55

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<210> 228

<211> 58

<212> PRT

<213> Homo sapiens

<400> 228

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Pro Glu Lys Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu Tyr
1           5           10           15
His Pro Asn Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu Tyr Cys
20           25           30

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<212> PRT

<213> Homo sapiens

<400> 231

Pro Lys Lys Asp Gly Ala Ala Thr Lys Val Asp Ala Ile Cys Thr Tyr
1 5 10 15

Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp
20 25 30

Glu Leu Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr
35 40 45

Gln Asp Arg Asp Ser Leu Tyr Asn Val Gly
50 55

<210> 232

<211> 58

<212> PRT

<213> Homo sapiens

<400> 232

Pro Glu Lys Asp Gly Ala Ala Thr Arg Val Asp Ala Val Cys Thr His
1 5 10 15

Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp
20 25 30

Lys Leu Ser Gln Leu Thr His Gly Ile Thr Glu Leu Gly Pro Tyr Thr
35 40 45

Leu Asp Arg His Ser Leu Tyr Val Asn Gly
50 55

<210> 233

<211> 58

<212> PRT

<213> Homo sapiens

<400> 233

Pro Glu Lys Asp Gly Val Ala Thr Arg Val Asp Ala Ile Cys Thr His
1 5 10 15

Arg Pro Asp Pro Lys Ile Pro Gly Leu Asp Arg Gln Gln Leu Tyr Trp
20 25 30

Glu Leu Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr
35 40 45

Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly
50 55

<210> 234

<211> 58

<212> PRT

<213> Homo sapiens

<400> 234

Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Ile His
1 5 10 15

His Leu Asp Pro Lys Ser Pro Gly Leu Asn Arg Glu Arg Leu Tyr Trp
20 25 30

Glu Leu Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr
35 40 45

Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly
50 55

<210> 235

<211> 58

<212> PRT

<213> Homo sapiens

<400> 235

Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His
1 5 10 15

Arg Leu Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp
20 25 30

Glu Leu Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr
35 40 45

Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly
50 55

<210> 236

<211> 58

<212> PRT

<213> Homo sapiens

<400> 236

Ser	Glu	Lys	Asp	Gly	Ala	Ala	Thr	Gly	Val	Asp	Ala	Ile	Cys	Thr	His
1				5					10					15	

Arg	Leu	Asp	Pro	Lys	Ser	Pro	Gly	Val	Asp	Arg	Glu	Gln	Leu	Tyr	Trp
			20					25					30		

Glu	Leu	Ser	Gln	Leu	Thr	Asn	Gly	Ile	Lys	Glu	Leu	Gly	Pro	Tyr	Thr
			35				40					45			

Leu	Asp	Arg	Asn	Ser	Leu	Tyr	Val	Asn	Gly
	50					55			

<210> 237

<211> 58

<212> PRT

<213> Homo sapiens

<400> 237

Ser	Glu	Lys	Asp	Gly	Ala	Ala	Thr	Gly	Val	Asp	Ala	Ile	Cys	Thr	His
1				5					10					15	

Arg	Val	Asp	Pro	Lys	Ser	Pro	Gly	Val	Asp	Arg	Glu	Gln	Leu	Tyr	Trp
			20					25					30		

Glu	Leu	Ser	Gln	Leu	Thr	Asn	Gly	Ile	Lys	Glu	Leu	Gly	Pro	Tyr	Thr
			35				40					45			

Leu	Asp	Arg	Asn	Ser	Leu	Tyr	Val	Asn	Gly
	50					55			

<210> 238

<211> 58

<212> PRT

<213> Homo sapiens

<400> 238

Ser	Glu	Lys	Asp	Gly	Ala	Ala	Thr	Gly	Val	Asp	Ala	Ile	Cys	Thr	His
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1 5 10 15
 His Leu Asn Pro Gln Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp
 20 25 30
 Gln Leu Ser Gln Met Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr
 35 40 45
 Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly
 50 55
 <210> 239
 <211> 58
 <212> PRT
 <213> Homo sapiens

 <400> 239
 Pro Glu Lys Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His
 1 5 10 15
 Arg Leu Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp
 20 25 30
 Glu Leu Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu
 35 40 45
 Leu Asp Arg Gly Ser Leu Tyr Val Asn Gly
 50 55
 <210> 240
 <211> 58
 <212> PRT
 <213> Homo sapiens

 <400> 240
 Pro Glu Lys Asn Gly Ala Ala Thr Gly Met Asp Ala Ile Cys Ser His
 1 5 10 15
 Arg Leu Asp Pro Lys Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp
 20 25 30
 Glu Leu Ser Gln Leu Thr His Gly Ile Lys Glu Leu Gly Pro Tyr Thr
 35 40 45
 Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly
 50 55

<210> 241

<211> 58

<212> PRT

<213> Homo sapiens

<400> 241

Pro Glu Lys Asn Gly Ala Ala Thr Gly Met Asp Ala Ile Cys Ser His
 1 5 10 15

Arg Leu Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp
 20 25 30

Glu Leu Ser Gln Leu Thr His Gly Ile Lys Glu Leu Gly Pro Tyr Thr
 35 40 45

Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly
 50 55

<210> 242

<211> 58

<212> PRT

<213> Homo sapiens

<400> 242

Pro Glu Lys His Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr Leu
 1 5 10 15

Arg Leu Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp
 20 25 30

Glu Leu Ser Gln Leu Thr Asn Ser Val Thr Glu Leu Gly Pro Tyr Thr
 35 40 45

Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly
 50 55

<210> 243

<211> 58

<212> PRT

<213> Homo sapiens

<400> 243

Pro Glu Lys His Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr Leu
1 5 10 15

Arg Leu Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp
20 25 30

Glu Leu Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr Thr
35 40 45

Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly
50 55

<210> 244

<211> 58

<212> PRT

<213> Homo sapiens

<400> 244

Pro Glu Lys His Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His
1 5 10 15

Arg Val Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp
20 25 30

Glu Leu Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr Thr
35 40 45

Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly
50 55

<210> 245

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<213> Homo sapiens

<400> 245

Pro Glu Lys Gln Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His
1 5 10 15

Arg Val Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp
20 25 30

Glu Leu Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr Thr
35 40 45

Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly

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<210> 246

<211> 58

<212> PRT

<213> Homo sapiens

<400> 246

Pro Glu Lys Gln Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His
 1 5 10 15

Arg Val Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp
 20 25 30

Glu Leu Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr Thr
 35 40 45

Leu Asp Arg Asp Ser Leu Tyr Val Asp Gly
 50 55

<210> 247

<211> 58

<212> PRT

<213> Homo sapiens

<400> 247

Pro Glu Lys Asp Lys Ala Ala Thr Arg Val Asp Ala Ile Cys Thr His
 1 5 10 15

His Pro Asp Pro Gln Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp
 20 25 30

Glu Leu Ser Gln Leu Thr His Gly Ile Thr Glu Leu Gly Pro Tyr Thr
 35 40 45

Leu Asp Arg Asp Ser Leu Tyr Val Asp Gly
 50 55

<210> 248

<211> 58

<212> PRT

<213> Homo sapiens

<400> 248

Ser Val Lys Asn Gly Ala Glu Thr Arg Val Asp Leu Leu Cys Thr Tyr
 1 5 10 15

Leu Gln Pro Leu Ser Gly Pro Gly Leu Pro Ile Lys Gln Val Phe His
 20 25 30

Glu Leu Ser Gln Gln Thr His Gly Ile Thr Arg Leu Gly Pro Tyr Ser
 35 40 45

Leu Asp Lys Asp Ser Leu Tyr Leu Asn Gly
 50 55

<210> 249

<211> 58

<212> PRT

<213> Homo sapiens

<400> 249

Pro Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Thr Thr Cys Thr Tyr
 1 5 10 15

His Pro Asp Pro Val Gly Pro Gly Leu Asp Ile Gln Gln Leu Tyr Trp
 20 25 30

Glu Leu Ser Gln Leu Thr His Gly Val Thr Gln Leu Gly Phe Tyr Val
 35 40 45

Leu Asp Arg Asp Ser Leu Phe Ile Asn Gly
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<210> 250

<211> 12

<212> PRT

<213> Homo sapiens

<400> 250

Phe Thr His Arg Ser Ser Met Pro Thr Thr Ser Thr
 1 5 10

<210> 251

<211> 12

<212> PRT

<213> Homo sapiens

<400> 251

Phe	Thr	His	Arg	Ser	Ser	Met	Pro	Thr	Thr	Ser	Ile
1				5					10		

<210> 252

<211> 12

<212> PRT

<213> Homo sapiens

<400> 252

Phe	Thr	His	Arg	Thr	Ser	Val	Pro	Thr	Ser	Ser	Thr
1				5					10		

<210> 253

<211> 12

<212> PRT

<213> Homo sapiens

<400> 253

Phe	Thr	His	Arg	Thr	Ser	Val	Pro	Thr	Thr	Ser	Thr
1				5					10		

<210> 254

<211> 12

<212> PRT

<213> Homo sapiens

<400> 254

Phe	Thr	His	Arg	Ser	Ser	Val	Pro	Thr	Thr	Ser	Ser
1				5					10		

<210> 255

<211> 12

<212> PRT

<213> Homo sapiens

<400> 255

Phe	Thr	His	Arg	Ser	Ser	Val	Ser	Thr	Thr	Ser	Thr
1				5					10		

<210> 256

<211> 12

<212> PRT

<213> Homo sapiens

<400> 256

Phe	Thr	His	Arg	Ser	Ser	Val	Ala	Pro	Thr	Ser	Thr
1				5					10		

<210> 257

<211> 12

<212> PRT

<213> Homo sapiens

<400> 257

Phe	Thr	His	Arg	Ser	Ser	Gly	Leu	Thr	Thr	Ser	Thr
1				5					10		

<210> 258

<211> 12

<212> PRT

<213> Homo sapiens

<400> 258

Phe	Thr	His	Arg	Ser	Phe	Gly	Leu	Thr	Thr	Ser	Thr
1				5					10		

<210> 259

<211> 12

<212> PRT

<213> Homo sapiens

<400> 259

Phe	Thr	His	Arg	Ser	Ser	Phe	Leu	Thr	Thr	Ser	Thr
1				5					10		

<210> 260

<211> 12

<212> PRT

<213> Homo sapiens

<400> 260

Phe	Thr	His	Arg	Asn	Phe	Val	Pro	Ile	Thr	Ser	Thr
1				5					10		

<210> 261

<211> 12

<212> PRT

<213> Homo sapiens

<400> 261

Phe	Thr	His	Arg	Ser	Ser	Val	Pro	Thr	Thr	Ser	Ile
1				5					10		

<210> 262

<211> 12

<212> PRT

<213> Homo sapiens

<400> 262

Phe	Thr	His	Gln	Ser	Ser	Val	Ser	Thr	Thr	Ser	Thr
1				5					10		

<210> 263

<211> 12

<212> PRT

<213> Homo sapiens

<400> 263

Phe Thr His Gln Thr Ser Ala Pro Asn Thr Ser Thr
1 5 10

<210> 264

<211> 12

<212> PRT

<213> Homo sapiens

<400> 264

Phe Thr His Gln Thr Phe Ala Pro Asn Thr Ser Thr
1 5 10

<210> 265

<211> 12

<212> PRT

<213> Homo sapiens

<400> 265

Phe Thr His Gln Asn Ser Val Pro Thr Thr Ser Thr
1 5 10

<210> 266

<211> 12

<212> PRT

<213> Homo sapiens

<400> 266

Phe Thr His Gln Ser Ser Met Thr Thr Thr Arg Thr

1 5 10
 <210> 267
 <211> 12
 <212> PRT
 <213> Homo sapiens

 <400> 267
 Phe Thr His Trp Ile Pro Val Pro Thr Ser Ser Thr
 1 5 10
 <210> 268
 <211> 12
 <212> PRT
 <213> Homo sapiens

 <400> 268
 Phe Thr His Trp Ser Pro Ile Pro Thr Thr Ser Thr
 1 5 10
 <210> 269
 <211> 12
 <212> PRT
 <213> Homo sapiens

 <400> 269
 Phe Thr His Trp Ser Ser Gly Leu Thr Thr Ser Thr
 1 5 10
 <210> 270
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 <213> Homo sapiens

 <400> 270

Phe	His	Pro	Arg	Ser	Ser	Val	Pro	Thr	Thr	Ser	Thr
1				5						10	

<210> 271

<211> 12

<212> PRT

<213> Homo sapiens

<400> 271

Phe	Asn	Pro	Arg	Ser	Ser	Val	Pro	Thr	Thr	Ser	Thr
1				5						10	

<210> 272

<211> 12

<212> PRT

<213> Homo sapiens

<400> 272

Phe	Asn	Pro	Trp	Ser	Ser	Val	Pro	Thr	Thr	Ser	Thr
1				5						10	

<210> 273

<211> 12

<212> PRT

<213> Homo sapiens

<400> 273

Phe	Thr	Gln	Arg	Ser	Ser	Val	Pro	Thr	Thr	Ser	Ile
1				5						10	

<210> 274

<211> 12

<212> PRT

<213> Homo sapiens

<400> 274

Phe Thr Gln Arg Ser Ser Val Pro Thr Thr Ser Thr
1 5 10

<210> 275

<211> 12

<212> PRT

<213> Homo sapiens

<400> 275

Phe Thr Gln Arg Ser Ser Val Pro Thr Thr Ser Val
1 5 10

<210> 276

<211> 12

<212> PRT

<213> Homo sapiens

<400> 276

Tyr Asn Glu Pro Gly Leu Asp Glu Pro Pro Thr Thr
1 5 10

<210> 277

<211> 12

<212> PRT

<213> Homo sapiens

<400> 277

Tyr Ala Pro Gln Asn Leu Ser Ile Arg Gly Glu Tyr
1 5 10

<210> 278

<211> 21

<212> PRT

<213> Homo sapiens

<400> 278

Pro Gly Thr Ser Thr Val Asp Val Gly Thr Ser Gly Thr Pro Ser Ser
 1 5 10 15

Ser Pro Ser Pro Thr
 20

<210> 279

<211> 23

<212> PRT

<213> Homo sapiens

<400> 279

Pro Gly Thr Ser Thr Val Asp Leu Arg Thr Ser Gly Thr Pro Ser Ser
 1 5 10 15

Leu Ser Ser Pro Thr Ile Met
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<210> 280

<211> 21

<212> PRT

<213> Homo sapiens

<400> 280

Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Phe Ser
 1 5 10 15

Leu Pro Ser Pro Ala
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<210> 281

<211> 20

<212> PRT

<213> Homo sapiens

<400> 281

Pro Gly Thr Ser Thr Val Asp Leu Gly Ser Gly Thr Pro Ser Ser Leu
 1 5 10 15

Pro Ser Pro Thr
20

<210> 282

<211> 20

<212> PRT

<213> Homo sapiens

<400> 282

Pro Gly Thr Ser Thr Val Asp Leu Gly Ser Gly Thr Pro Ser Leu Pro
1 5 10 15

Ser Ser Pro Thr
20

<210> 283

<211> 21

<212> PRT

<213> Homo sapiens

<400> 283

Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Ser
1 5 10 15

Leu Pro Ser Pro Thr
20

<210> 284

<211> 21

<212> PRT

<213> Homo sapiens

<400> 284

Pro Gly Thr Pro Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Val Ser
1 5 10 15

Lys Pro Gly Pro Ser
20

<210> 285

<211> 21

<212> PRT

<213> Homo sapiens

<400> 285

Pro Trp Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Pro
1 5 10 15

Val Pro Ser Pro Thr
20

<210> 286

<211> 21

<212> PRT

<213> Homo sapiens

<400> 286

Pro Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr Gly Thr Pro Ser Ser
1 5 10 15

Phe Pro Gly His Thr
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<210> 287

<211> 21

<212> PRT

<213> Homo sapiens

<400> 287

Pro Gly Thr Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Ser
1 5 10 15

Leu Pro Gly His Thr
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<210> 288

<211> 21

<212> PRT

<213> Homo sapiens

<400> 288

Pro Gly Thr Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Pro
1 5 10 15

Leu Pro Gly His Thr
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<210> 289

<211> 21

<212> PRT

<213> Homo sapiens

<400> 289

Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser Arg Thr Pro Ala Ser
1 5 10 15

Leu Ser Gly Pro Thr
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<210> 290

<211> 21

<212> PRT

<213> Homo sapiens

<400> 290

Pro Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly Thr Pro Ala Ser
1 5 10 15

Leu Pro Gly His Thr
20

<210> 291

<211> 21

<212> PRT

<213> Homo sapiens

<400> 291

Pro Gly Thr Ser Ala Val His Leu Glu Thr Thr Gly Thr Pro Ser Ser
 1 5 10 15

Phe Pro Gly His Thr
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<210> 292

<211> 21

<212> PRT

<213> Homo sapiens

<400> 292

Pro Gly Thr Ser Thr Val His Leu Gly Thr Ser Glu Thr Pro Ser Ser
 1 5 10 15

Leu Pro Arg Pro Ile
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<210> 293

<211> 21

<212> PRT

<213> Homo sapiens

<400> 293

Pro Gly Thr Ser Ile Val Asn Leu Gly Thr Ser Gly Ile Pro Pro Ser
 1 5 10 15

Leu Pro Glu Thr Thr
 20

<210> 294

<211> 21

<212> PRT

<213> Homo sapiens

<400> 294

Pro Gly Thr Phe Thr Val Gln Pro Glu Thr Ser Glu Thr Pro Ser Ser
 1 5 10 15

Leu Pro Gly Pro Thr

20

<210> 295

<211> 21

<212> PRT

<213> Homo sapiens

<400> 295

Pro	Gly	Thr	Pro	Thr	Val	Asp	Leu	Gly	Thr	Ser	Gly	Thr	Pro	Val	Ser
1				5					10					15	

Lys	Pro	Gly	Pro	Ser
			20	

<210> 296

<211> 21

<212> PRT

<213> Homo sapiens

<400> 296

Pro	Gly	Thr	Pro	Thr	Val	Tyr	Leu	Gly	Ala	Ser	Lys	Thr	Pro	Ala	Ser
1				5					10					15	

Ile	Phe	Gly	Pro	Ser
			20	

<210> 297

<211> 16

<212> PRT

<213> Homo sapiens

<400> 297

Pro	Lys	Pro	Ala	Thr	Thr	Phe	Leu	Pro	Pro	Leu	Ser	Glu	Ala	Thr	Thr
1				5					10					15	

<210> 298

<211> 21

<212> PRT

<213> Homo sapiens

<400> 298

Gln Ile Asn Phe His Ile Val Asn Trp Asn Leu Ser Asn Pro Asp Pro
 1 5 10 15
 Thr Ser Ser Glu Tyr
 20

<210> 299

<211> 1794

<212> PRT

<213> Homo sapiens

<400> 299

Met Glu His Ile Thr Lys Ile Pro Asn Glu Ala Ala His Arg Gly Thr
 1 5 10 15
 Ile Arg Pro Val Lys Gly Pro Gln Thr Ser Thr Ser Pro Ala Ser Pro
 20 25 30
 Lys Gly Leu His Thr Gly Gly Thr Lys Arg Met Glu Thr Thr Thr Thr
 35 40 45
 Ala Leu Lys Thr Thr Thr Thr Ala Leu Lys Thr Thr Ser Arg Ala Thr
 50 55 60
 Leu Thr Thr Ser Val Tyr Thr Pro Thr Leu Gly Thr Leu Thr Pro Leu
 65 70 75 80
 Asn Ala Ser Arg Gln Met Ala Ser Thr Ile Leu Thr Glu Met Met Ile
 85 90 95
 Thr Thr Pro Tyr Val Phe Pro Asp Val Pro Glu Thr Thr Ser Ser Leu
 100 105 110
 Ala Thr Ser Leu Gly Ala Glu Thr Ser Thr Ala Leu Pro Arg Thr Thr
 115 120 125
 Pro Ser Val Leu Asn Arg Glu Ser Glu Thr Thr Ala Ser Leu Val Ser
 130 135 140
 Arg Ser Gly Ala Glu Arg Ser Pro Val Ile Gln Thr Leu Asp Val Ser
 145 150 155 160
 Ser Ser Glu Pro Asp Thr Thr Ala Ser Trp Val Ile His Pro Ala Glu
 165 170 175
 Thr Ile Pro Thr Val Ser Lys Thr Thr Pro Asn Phe Phe His Ser Glu
 180 185 190

Leu Asp Thr Val Ser Ser Thr Ala Thr Ser His Gly Ala Asp Val Ser
 195 200 205
 Ser Ala Ile Pro Thr Asn Ile Ser Pro Ser Glu Leu Asp Ala Leu Thr
 210 215 220
 Pro Leu Val Thr Ile Ser Gly Thr Asp Thr Ser Thr Thr Phe Pro Thr
 225 230 235 240
 Leu Thr Lys Ser Pro His Glu Thr Glu Thr Arg Thr Thr Trp Leu Thr
 245 250 255
 His Pro Ala Glu Thr Ser Ser Thr Ile Pro Arg Thr Ile Pro Asn Phe
 260 265 270
 Ser His His Glu Ser Asp Ala Thr Pro Ser Ile Ala Thr Ser Pro Gly
 275 280 285
 Ala Glu Thr Ser Ser Ala Ile Pro Ile Met Thr Val Ser Pro Gly Ala
 290 295 300
 Glu Asp Leu Val Thr Ser Gln Val Thr Ser Ser Gly Thr Asp Arg Asn
 305 310 315 320
 Met Thr Ile Pro Thr Leu Thr Leu Ser Pro Gly Glu Pro Lys Thr Ile
 325 330 335
 Ala Ser Leu Val Thr His Pro Glu Ala Gln Thr Ser Ser Ala Ile Pro
 340 345 350
 Thr Ser Thr Ile Ser Pro Ala Val Ser Arg Leu Val Thr Ser Met Val
 355 360 365
 Thr Ser Leu Ala Ala Lys Thr Ser Thr Thr Asn Arg Ala Leu Thr Asn
 370 375 380
 Ser Pro Gly Glu Pro Ala Thr Thr Val Ser Leu Val Thr His Pro Ala
 385 390 395 400
 Gln Thr Ser Pro Thr Val Pro Trp Thr Thr Ser Ile Phe Phe His Ser
 405 410 415
 Lys Ser Asp Thr Thr Pro Ser Met Thr Thr Ser His Gly Ala Glu Ser
 420 425 430
 Ser Ser Ala Val Pro Thr Pro Thr Val Ser Thr Glu Val Pro Gly Val
 435 440 445
 Val Thr Pro Leu Val Thr Ser Ser Arg Ala Val Ile Ser Thr Thr Ile
 450 455 460
 Pro Ile Leu Thr Leu Ser Pro Gly Glu Pro Glu Thr Thr Pro Ser Met
 465 470 475 480
 Ala Thr Ser His Gly Glu Glu Ala Ser Ser Ala Ile Pro Thr Pro Thr
 485 490 495
 Val Ser Pro Gly Val Pro Gly Val Val Thr Ser Leu Val Thr Ser Ser

500	505	510
Arg Ala Val Thr Ser Thr Thr Ile Pro Ile Leu Thr Phe Ser Leu Gly 515 520 525		
Glu Pro Glu Thr Thr Pro Ser Met Ala Thr Ser His Gly Thr Glu Ala 530 535 540		
Gly Ser Ala Val Pro Thr Val Leu Pro Glu Val Pro Gly Met Val Thr 545 550 555 560		
Ser Leu Val Ala Ser Ser Arg Ala Val Thr Ser Thr Thr Leu Pro Thr 565 570 575		
Leu Thr Leu Ser Pro Gly Glu Pro Glu Thr Thr Pro Ser Met Ala Thr 580 585 590		
Ser His Gly Ala Glu Ala Ser Ser Thr Val Pro Thr Val Ser Pro Glu 595 600 605		
Val Pro Gly Val Val Thr Ser Leu Val Thr Ser Ser Ser Gly Val Asn 610 615 620		
Ser Thr Ser Ile Pro Thr Leu Ile Leu Ser Pro Gly Glu Leu Glu Thr 625 630 635 640		
Thr Pro Ser Met Ala Thr Ser His Gly Ala Glu Ala Ser Ser Ala Val 645 650 655		
Pro Thr Pro Thr Val Ser Pro Gly Val Ser Gly Val Val Thr Pro Leu 660 665 670		
Val Thr Ser Ser Arg Ala Val Thr Ser Thr Thr Ile Pro Ile Leu Thr 675 680 685		
Leu Ser Ser Ser Glu Pro Glu Thr Thr Pro Ser Met Ala Thr Ser His 690 695 700		
Gly Val Glu Ala Ser Ser Ala Val Leu Thr Val Ser Pro Glu Val Pro 705 710 715 720		
Gly Met Val Thr Ser Leu Val Thr Ser Ser Arg Ala Val Thr Ser Thr 725 730 735		
Thr Ile Pro Thr Leu Thr Ile Ser Ser Asp Glu Pro Glu Thr Thr Thr 740 745 750		
Ser Leu Val Thr His Ser Glu Ala Lys Met Ile Ser Ala Ile Pro Thr 755 760 765		
Leu Ala Val Ser Pro Thr Val Gln Gly Leu Val Thr Ser Leu Val Thr 770 775 780		
Ser Ser Gly Ser Glu Thr Ser Ala Phe Ser Asn Leu Thr Val Ala Ser 785 790 795 800		
Ser Gln Pro Glu Thr Ile Asp Ser Trp Val Ala His Pro Gly Thr Glu 805 810 815		

Ala Ser Ser Val Val Pro Thr Leu Thr Val Ser Thr Gly Glu Pro Phe
 820 825 830
 Thr Asn Ile Ser Leu Val Thr His Pro Ala Glu Ser Ser Ser Thr Leu
 835 840 845
 Pro Arg Thr Thr Ser Arg Phe Ser His Ser Glu Leu Asp Thr Met Pro
 850 855 860
 Ser Thr Val Thr Ser Pro Glu Ala Glu Ser Ser Ser Ala Ile Ser Thr
 865 870 875 880
 Thr Ile Ser Pro Gly Ile Pro Gly Val Leu Thr Ser Leu Val Thr Ser
 885 890 895
 Ser Gly Arg Asp Ile Ser Ala Thr Phe Pro Thr Val Pro Glu Ser Pro
 900 905 910
 His Glu Ser Glu Ala Thr Ala Ser Trp Val Thr His Pro Ala Val Thr
 915 920 925
 Ser Thr Thr Val Pro Arg Thr Thr Pro Asn Tyr Ser His Ser Glu Pro
 930 935 940
 Asp Thr Thr Pro Ser Ile Ala Thr Ser Pro Gly Ala Glu Ala Thr Ser
 945 950 955 960
 Asp Phe Pro Thr Ile Thr Val Ser Pro Asp Val Pro Asp Met Val Thr
 965 970 975
 Ser Gln Val Thr Ser Ser Gly Thr Asp Thr Ser Ile Thr Ile Pro Thr
 980 985 990
 Leu Thr Leu Ser Ser Gly Glu Pro Glu Thr Thr Thr Ser Phe Ile Thr
 995 1000 1005
 Tyr Ser Glu Thr His Thr Ser Ser Ala Ile Pro Thr Leu Pro Val
 1010 1015 1020
 Ser Pro Gly Ala Ser Lys Met Leu Thr Ser Leu Val Ile Ser Ser
 1025 1030 1035
 Gly Thr Asp Ser Thr Thr Thr Phe Pro Thr Leu Thr Glu Thr Pro
 1040 1045 1050
 Tyr Glu Pro Glu Thr Thr Ala Ile Gln Leu Ile His Pro Ala Glu
 1055 1060 1065
 Thr Asn Thr Met Val Pro Arg Thr Thr Pro Lys Phe Ser His Ser
 1070 1075 1080
 Lys Ser Asp Thr Thr Leu Pro Val Ala Ile Thr Ser Pro Gly Pro
 1085 1090 1095
 Glu Ala Ser Ser Ala Val Ser Thr Thr Thr Ile Ser Pro Asp Met
 1100 1105 1110
 Ser Asp Leu Val Thr Ser Leu Val Pro Ser Ser Gly Thr Asp Thr
 1115 1120 1125

Ser Thr Thr Phe Pro Thr Leu Ser Glu Thr Pro Tyr Glu Pro Glu
 1130 1135 1140
 Thr Thr Ala Thr Trp Leu Thr His Pro Ala Glu Thr Ser Thr Thr
 1145 1150 1155
 Val Ser Gly Thr Ile Pro Asn Phe Ser His Arg Gly Ser Asp Thr
 1160 1165 1170
 Ala Pro Ser Met Val Thr Ser Pro Gly Val Asp Thr Arg Ser Gly
 1175 1180 1185
 Val Pro Thr Thr Thr Ile Pro Pro Ser Ile Pro Gly Val Val Thr
 1190 1195 1200
 Ser Gln Val Thr Ser Ser Ala Thr Asp Thr Ser Thr Ala Ile Pro
 1205 1210 1215
 Thr Leu Thr Pro Ser Pro Gly Glu Pro Glu Thr Thr Ala Ser Ser
 1220 1225 1230
 Ala Thr His Pro Gly Thr Gln Thr Gly Phe Thr Val Pro Ile Arg
 1235 1240 1245
 Thr Val Pro Ser Ser Glu Pro Asp Thr Met Ala Ser Trp Val Thr
 1250 1255 1260
 His Pro Pro Gln Thr Ser Thr Pro Val Ser Arg Thr Thr Ser Ser
 1265 1270 1275
 Phe Ser His Ser Ser Pro Asp Ala Thr Pro Val Met Ala Thr Ser
 1280 1285 1290
 Pro Arg Thr Glu Ala Ser Ser Ala Val Leu Thr Thr Ile Ser Pro
 1295 1300 1305
 Gly Ala Pro Glu Met Val Thr Ser Gln Ile Thr Ser Ser Gly Ala
 1310 1315 1320
 Ala Thr Ser Thr Thr Val Pro Thr Leu Thr His Ser Pro Gly Met
 1325 1330 1335
 Pro Glu Thr Thr Ala Leu Leu Ser Thr His Pro Arg Thr Glu Thr
 1340 1345 1350
 Ser Lys Thr Phe Pro Ala Ser Thr Val Phe Pro Gln Val Ser Glu
 1355 1360 1365
 Thr Thr Ala Ser Leu Thr Ile Arg Pro Gly Ala Glu Thr Ser Thr
 1370 1375 1380
 Ala Leu Pro Thr Gln Thr Thr Ser Ser Leu Phe Thr Leu Leu Val
 1385 1390 1395
 Thr Gly Thr Ser Arg Val Asp Leu Ser Pro Thr Ala Ser Pro Gly
 1400 1405 1410
 Val Ser Ala Lys Thr Ala Pro Leu Ser Thr His Pro Gly Thr Glu

1415		1420		1425
Thr Ser	Thr Met Ile Pro	Thr Ser Thr Leu Ser	Leu Gly Leu Leu	
1430		1435	1440	
Glu Thr	Thr Gly Leu Leu Ala	Thr Ser Ser Ser	Ala Glu Thr Ser	
1445		1450	1455	
Thr Ser	Thr Leu Thr Leu Thr	Val Ser Pro Ala Val	Ser Gly Leu	
1460		1465	1470	
Ser Ser	Ala Ser Ile Thr Thr	Asp Lys Pro Gln Thr	Val Thr Ser	
1475		1480	1485	
Trp Asn	Thr Glu Thr Ser Pro	Ser Val Thr Ser Val	Gly Pro Pro	
1490		1495	1500	
Glu Phe	Ser Arg Thr Val Thr	Gly Thr Thr Met Thr	Leu Ile Pro	
1505		1510	1515	
Ser Glu	Met Pro Thr Pro Pro	Lys Thr Ser His Gly	Glu Gly Val	
1520		1525	1530	
Ser Pro	Thr Thr Ile Leu Arg	Thr Thr Met Val Glu	Ala Thr Asn	
1535		1540	1545	
Leu Ala	Thr Thr Gly Ser Ser	Pro Thr Val Ala Lys	Thr Thr Thr	
1550		1555	1560	
Thr Phe	Asn Thr Leu Ala Gly	Ser Leu Phe Thr Pro	Leu Thr Thr	
1565		1570	1575	
Pro Gly	Met Ser Thr Leu Ala	Ser Glu Ser Val Thr	Ser Arg Thr	
1580		1585	1590	
Ser Tyr	Asn His Arg Ser Trp	Ile Ser Thr Thr Ser	Ser Tyr Asn	
1595		1600	1605	
Arg Arg	Tyr Trp Thr Pro Ala	Thr Ser Thr Pro Val	Thr Ser Thr	
1610		1615	1620	
Phe Ser	Pro Gly Ile Ser Thr	Ser Ser Ile Pro Ser	Ser Thr Ala	
1625		1630	1635	
Ala Thr	Val Pro Phe Met Val	Pro Phe Thr Leu Asn	Phe Thr Ile	
1640		1645	1650	
Thr Asn	Leu Gln Tyr Glu Glu	Asp Met Arg His Pro	Gly Ser Arg	
1655		1660	1665	
Lys Phe	Asn Ala Thr Glu Arg	Glu Leu Gln Gly Leu	Leu Lys Pro	
1670		1675	1680	
Leu Phe	Arg Asn Ser Ser Leu	Glu Tyr Leu Tyr Ser	Gly Cys Arg	
1685		1690	1695	
Leu Ala	Ser Leu Arg Pro Glu	Lys Asp Ser Ser Ala	Met Ala Val	
1700		1705	1710	

Asp Ala Ile Cys Thr His Arg Pro Asp Pro Glu Asp Leu Gly Leu
 1715 1720 1725
 Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Asn Leu Thr Asn Gly
 1730 1735 1740
 Ile Gln Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr
 1745 1750 1755
 Val Asn Gly Phe Thr His Arg Ser Ser Met Pro Thr Thr Ser Thr
 1760 1765 1770
 Pro Gly Thr Ser Thr Val Asp Val Gly Thr Ser Gly Thr Pro Ser
 1775 1780 1785
 Ser Ser Pro Ser Pro Thr
 1790

<210> 300

<211> 284

<212> PRT

<213> Homo sapiens

<400> 300

Ile Thr Leu Leu Arg Asp Ile Gln Asp Lys Val Thr Thr Leu Tyr Lys
 1 5 10 15
 Gly Ser Gln Leu His Asp Thr Phe Arg Phe Cys Leu Val Thr Asn Leu
 20 25 30
 Thr Met Asp Ser Val Leu Val Thr Val Lys Ala Leu Phe Ser Ser Asn
 35 40 45
 Leu Asp Pro Ser Leu Val Glu Gln Val Phe Leu Asp Lys Thr Leu Asn
 50 55 60
 Ala Ser Phe His Trp Leu Gly Ser Thr Tyr Gln Leu Val Asp Ile His
 65 70 75 80
 Val Thr Glu Met Glu Ser Ser Val Tyr Gln Pro Thr Ser Ser Ser Ser
 85 90 95
 Thr Gln His Phe Tyr Leu Asn Phe Thr Ile Thr Asn Leu Pro Tyr Ser
 100 105 110
 Gln Asp Lys Ala Gln Pro Gly Thr Thr Asn Tyr Gln Arg Asn Lys Arg
 115 120 125
 Asn Ile Glu Asp Ala Leu Asn Gln Leu Phe Arg Asn Ser Ser Ile Lys
 130 135 140
 Ser Tyr Phe Ser Asp Cys Gln Val Ser Thr Phe Arg Ser Val Pro Asn
 145 150 155 160

Arg His His Thr Gly Val Asp Ser Leu Cys Asn Phe Ser Pro Leu Ala
165 170 175

Arg Arg Val Asp Arg Val Ala Ile Tyr Glu Glu Phe Leu Arg Met Thr
180 185 190

Arg Asn Gly Thr Gln Leu Gln Asn Phe Thr Leu Asp Arg Ser Ser Val
195 200 205

Leu Val Asp Gly Tyr Ser Pro Asn Arg Asn Glu Pro Leu Thr Gly Asn
210 215 220

Ser Asp Leu Pro Phe Trp Ala Val Ile Leu Ile Gly Leu Ala Gly Leu
225 230 235 240

Leu Gly Leu Ile Thr Cys Leu Ile Cys Gly Val Leu Val Thr Thr Arg
245 250 255

Arg Arg Lys Lys Glu Gly Glu Tyr Asn Val Gln Gln Gln Cys Pro Gly
260 265 270

Tyr Tyr Gln Ser His Leu Asp Leu Glu Asp Leu Gln
275 280

<210> 301

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic primer

<400> 301
gtctctatgt caatggtttc accc

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<210> 302

<211> 21

<212> DNA

<213> Artificial Sequence

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<223> Synthetic primer

<400> 302
tagctgctct ctgtccagtc c

21

<210> 303
 <211> 22
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetic primer
 <400> 303
 ggacaaggtc accacactct ac

22

<210> 304
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
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 <400> 304
 gcagatcctc caggtctagg tgtg

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<210> 305
 <211> 24
 <212> DNA
 <213> Artificial Sequence

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 <400> 305
 gtctctatgt caatggtttc accc

24

<210> 306

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic primer

<400> 306

tagctgctct ctgtccagtc c

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